# **Human Secreted Proteins**

This application is a continuation-in-part of, and claims benefit under 35 U.S.C. [1] § 119(e) based on copending U.S. Provisional Application No. 60/278,650 filed on March 27, 2001. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending U.S. Utility Application No. 09/833,245, filed on April 12, 2001, and PCT International Application Serial No. US01/11988, filed on April 12, 2001. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06043, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/167,061, filed on November 23, 1999, and U.S. Provisional Application No. 60/124,146, filed on March 12, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. U\$00/06012, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/166,989, filed on November 23, 1999, and U.S. Provisional Application No. 60/124,093, filed on March 12, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06058, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/168,654, filed on December 3, 1999, and U.S. Provisional Application No. 60/124,145, filed on March 12, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06044, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/168,661, filed on December 3, 1999, and U.S. Provisional Application No. 60/124,099, filed on March 12, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT\International Application Serial No. US00/06059, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/168,622, filed on December 3, 1999, and U.S. Provisional Application No. 60/124,096, filed on March 12, 1999. This application

is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06042, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/168,663, filed on December 3, 1999, and U.S. Provisional Application No. 60/124,143, filed on March 12, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06014, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/168,665, filed on December 3, 1999, and U.S. Provisional Application No. 60/138,598, filed on June 11, 1999, and U.S. Provisional Application No. 60/124,095, filed on March 12, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06013, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 1/19(e) based on U.S. Provisional Application No. 60/168,662, filed on December 3, 1999, and U.S. Provisional Application No. 60/138,626, filed on June 11,1999, and U.S. Provisional Application No. 60/125,360, filed on March 19, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06049, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/168,667, filed on December 3, 1999, and U.S. Provisional Application No. 60/138,574, filed on June 11, 1999, and U.S. Provisional Application No. 60/124,144, filed on March 12, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of dopending PCT International Application Serial No. US00/06057, filed on March 9, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/168,666, filed on December 3, 1999, and U.S. Provisional Application No. 60/134,597, filed on June 11, 1999, and U.S. Provisional Application No. 60/124,142, filed on March 12, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00 (06824, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/168,664, filed on December 3, 1999, and U.S. Provisional Application No. 60/125,359, filed on March 19, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06765, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/169,906, filed on December 10, 1999, and

U.S. Provisional Application No. 60/126,051, filed on March 23, 1999. This application is also a continuation-in part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06792, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/169,980, filed on December 10, 1999, and U.S. Provisional Application No. 60/125,362, filed on March 19, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06830, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provistonal Application No. 60/169,910, filed on December 10, 1999, and U.S. Provisional Application No. 60/125,361, filed on March 19, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06782, filed on March 16, 2000, which claims benefit uhder 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/169,936, filed on December 10, 1999, and U.S. Provisional Application No. 60/125,812, filed on March 23, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06822, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/169,916, filed on December 10, 1999, and US. Provisional Application No. 60/126,054, filed on March 23, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06791, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/169,946, filed on December 10, 1999, and U.S. Provisional Application No. 60/123,815, filed on March 23, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No.\US00/06828, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/169,616, filed on December 8, 1999, and U.S. Provisional Application No. 60/125,358, filed on March 19, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06823, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/169,623, filed on December 8, 1999, and U.S. Provisional Application No. 60/125,364, filed on March 19, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/06781, filed on March 16, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/169,617, filed on December 8, 1999, and U.S. Provisional Application No. 60/125,363, filed on March 19, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07505, filed on March 22, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/172,410, filed on December 17, 1999, and U.S. Provisional Application No. 60/126,502, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07440, filed on March 22, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/172,409, filed on December 17, 1999, and U.S. Provisional Application No. 60/126,503, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under §5 U.S.C. § 120 of copending PCT International Application Serial No. US00/07506, filed on March 22, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/172,412, filed on December 17, 1999, and U.S. Provisional Application No. 60/126,505, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07507, filed on March 22, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/172,408, filed on December 17, 1999, and U.S. Provisional Application No. 60/126,594, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/172,413, filed on December 17, 1999, and U.S. Provisional Application No. 60/126,511, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07525, filed on March 22, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/171,549, filed on December 22, 1999, and U.S. Provisional Application No. 60/126,595, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07534, filed on March 22, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No.

60/171,504, filed on December 22, 1999, and U.S. Provisional Application No. 60/126,598, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07483, filed on March 22, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/171,552, filed on December 22, 1999, and U.S. Provisional Application No. 60/126,596, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07526, filed on March 22, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60(171,550, filed on December 22, 1999, and U.S. Provisional Application No. 60/126,600, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07527, filed on March 22, 2000, which claims benefit under 35 U.S.C. § 19(e) based on U.S. Provisional Application No. 60/171,551, filed on December 22, 1999, and U.S. Provisional Application No. 60/126,501, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07661, filed on March 23 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/174,847, filed on January 7, 2000, and U.S. Provisional Application No. 60/126\504, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. U\$00/07579, filed on March 23, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/174,853, filed on January 7, 2000, and U.S. Provisional Application No. 60/126,509, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07723, filed on March 23, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/242,710, filed on October 25, 2000, and U.S. Provisional Application No. 60/174,850, filed on January 7, 2000, and U.S. Provisional Application No. 60/126,506, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07724, filed on March 23, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/174,850, filed on January 7, 2000, and U.S. Provisional Application No. 60/126,510, filed on March \$6, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/14929, filed on June 1, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/174,851, filed on January 7, 2000, and U.S. Provisional Application No. 60/138,573, filed on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07722, filed on March 23, 2000, which claims benefit under 35\U.S.C. § 119(e) based on U.S. Provisional Application No. 60/174,871, filed on January 7, 2000, and U.S. Provisional Application No. 60/126,508, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit § 120 of copending PCT International Application Serial No. under 35 U.S.C. US00/07578, filed on March 23, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/174,872, filed on January 7, 2000, and U.S. Provisional Application No. 60/126,507, filed on March 26, 1999. This application is also a continuation-in-part of,\and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07726, filed on March 23, 2000, which claims benefit under 35 U.S.C. § 1\( 9(e) \) based on U.S. Provisional Application No. 60/174,877, filed on January 7, 2000, and U.S. Provisional Application No. 60/126,597, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/07677, filed on March 23\, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/176,064, filed on January 14, 2000, and U.S. Provisional Application No. \$0/154,373, filed on September 17, 1999, and U.S. Provisional Application No. 60/126,601, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. U\$00/07725, filed on March 23, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/176,063, filed on January 14, 2000, and U.S. Provisional Application No. 60/126,602, filed on March 26, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/09070, filed on April 6, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/176,052, filed on January 14, 2000, and U.S. Provisional Application No. 60/128,695, filed on April 9, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT

International Application Serial No. US00/08982, filed on April 6, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/176,069, filed on January 14, 2000, and U.S. Provisional Application No. 60/128,696, filed on April 9, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/08983, filed on April 6, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/176,068, filed on January 14, 2000, and U.S. Provisional Application No. 60/128,703, filed on April 9, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/09067, filed on April 6, 2000, which claims benefit under 35 U.S.¢. § 119(e) based on U.S. Provisional Application No. 60/176,929, filed on January 20, 2000, and U.S. Provisional Application No. 60/128,697, filed on April 9, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/09066, filed on April 6, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/176,926, and U.S. Provisional Application No. 60/128,698, filed on April 9, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.Q. § 120 of copending PCT International Application Serial No. US00/09068, filed on April 6, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/177,050, filed on January 20, 2000, and U.S. Provisional Application No. 60/128,699, filed on April 9, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial Nd US00/08981, filed on April 6, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/177,166, filed on January 20, 2000, and U.S.\Provisional Application No. 60/128,701, filed on April 9, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/08980, filed on April 6, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/176,930, filed on January 20, 2000, and U.S. Provisional Application No. 60/128,700, filed on April 9, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/09071, filed on April 6, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.\$. Provisional Application No. 60/176,931, filed on January 20, 2000, and U.S. Provisional Application No. 60/128,694, filed on

April 9, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/09069, filed on April 6, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/177,049, filed on January 20, 2000, and U.S. Provisional Application No. 60/128,702, filed on April 9, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/15136, filed on June 1, 2000, which claims benefit under 35 U.S.C\ § 119(e) based on U.S. Provisional Application No. 60/138,629, filed on June 11, 1999.\ This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/14926, filed on June 1, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/138,628, filed on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/14963, filed on June 1, 2000, which claims benefit under 35 U.S\C. § 119(e) based on U.S. Provisional Application No. 60/138,631, filed on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/15135, filed on June 1, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/138,632, filed on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/14934, filed on June 1, 2000, which claims benefit under 35 U.S.C.\§ 119(e) based on U.S. Provisional Application No. 60/138,599, filed on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/14933, filed on June 1, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/138,572, filed on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/15137, filed on June 1, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/138,625, filed on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/14928, filed on June 1, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60(138,633, filed on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of

copending PCT International Application Serial No. US00/14973, filed on June 1, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/,138,630, filed on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/14964, filed on June 1, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/138,627, filed on June 11, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/26376, filed on September 26, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/155,808, filed on September 27, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/26371, filed on September 26, 2000, which claims benefit under 35 U.S.C.\§ 119(e) based on U.S. Provisional Application No. 60/155,804, filed on September 2λ, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/26324, filed on September 26, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/155,807, filed on September 27, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/26323, filed on September 26, 2000, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/155,805, filed on September 27, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US00/26337, filed on September 26, 2000, which claims benefit under 35 U.S.Q § 119(e) based on U.S. Provisional Application No. 60/155,806, filed on September 27, 1999. This application is also a continuation-in-part of, and claims benefit under 35 U.S.C. § 120 of copending PCT International Application Serial No. US01/13318, filed on April 27, 2001, which claims benefit under 35 U.S.C. § 119(e) based on U.S. Provisional Application No. 60/212,142, filed on June 16, 2000, and U.S. Provisional Application No. 60/201,194, filed on May 2, 2000. Each of the above referenced PCT applications were published in the English language. Each of the above referenced priority applications are hereby incorporated by reference in their entireties.

## Field of the Invention

The present invention relates to novel proteins. More specifically, isolated nucleic acid molecules are provided encoding novel polypeptides. Novel polypeptides and antibodies that bind to these polypeptides are provided. Also provided are vectors, host cells, and recombinant and synthetic methods for producing human polynucleotides and/or polypeptides, and antibodies. The invention further relates to diagnostic and therapeutic methods useful for diagnosing, treating, preventing and/or prognosing disorders related to these novel polypeptides. The invention further relates to screening methods for identifying agonists and antagonists of polynucleotides and polypeptides of the invention. The present invention further relates to methods and/or compositions for inhibiting or enhancing the production and function of the polypeptides of the present invention.

## Background of the Invention

- [3] Unlike bacterium, which exist as a single compartment surrounded by a membrane, human cells and other eukaryotes are subdivided by membranes into many functionally distinct compartments. Each membrane-bounded compartment, or organelle, contains different proteins essential for the function of the organelle. The cell uses "sorting signals," which are amino acid motifs located within the protein, to target proteins to particular cellular organelles.
- One type of sorting signal, called a signal sequence, a signal peptide, or a leader sequence, directs a class of proteins to an organelle called the endoplasmic reticulum (ER). The ER separates the membrane-bounded proteins from all other types of proteins. Once localized to the ER, both groups of proteins can be further directed to another organelle called the Golgi apparatus. Here, the Golgi distributes the proteins to vesicles, including secretory vesicles, the cell membrane, lysosomes, and the other organelles.
- [5] Proteins targeted to the ER by a signal sequence can be released into the extracellular space as a secreted protein. For example, vesicles containing secreted proteins can fuse with the cell membrane and release their contents into the extracellular space a process called exocytosis. Exocytosis can occur constitutively or after receipt of a triggering signal. In the latter case, the proteins are stored in secretory vesicles (or

secretory granules) until exocytosis is triggered. Similarly, proteins residing on the cell membrane can also be secreted into the extracellular space by proteolytic cleavage of a "linker" holding the protein to the membrane.

Thus there exists a clear need for identifying and using novel secreted polynucleotides and polypeptides. Identification and sequencing of human genes is a major goal of modern scientific research. For example, by identifying genes and determining their sequences, scientists have been able to make large quantities of valuable human "gene products." These include human insulin, interferon, Factor VIII, tumor necrosis factor, human growth hormone, tissue plasminogen activator, and numerous other compounds. Additionally, knowledge of gene sequences can provide the key to treatment or cure of genetic diseases (such as muscular dystrophy and cystic fibrosis).

## Summary of the Invention

The present invention relates to novel secreted proteins. More specifically, isolated nucleic acid molecules are provided encoding novel secreted polypeptides. Novel polypeptides and antibodies that bind to these polypeptides are provided. Also provided are vectors, host cells, and recombinant and synthetic methods for producing human polynucleotides and/or polypeptides, and antibodies. The invention further relates to diagnostic and therapeutic methods useful for diagnosing, treating, preventing and/or prognosing disorders related to these novel polypeptides. The invention further relates to screening methods for identifying agonists and antagonists of polynucleotides and polypeptides of the invention. The present invention further relates to methods and/or compositions for inhibiting or enhancing the production and function of the polypeptides of the present invention.

#### **Detailed Description**

### Polynucleotides and Polypeptides

## Description of Table 1A

[8] Table 1A summarizes information concerning certain polypnucleotides and polypeptides of the invention. The first column provides the gene number in the

application for each clone identifier. The second column provides a unique clone identifier, "Clone ID NO:Z", for a cDNA clone related to each contig sequence disclosed in Table 1A. Third column, the cDNA Clones identified in the second column were deposited as indicated in the third column (i.e. by ATCC Deposit Number and deposit date). Some of the deposits contain multiple different clones corresponding to the same gene. In the fourth column, "Vector" refers to the type of vector contained in the corresponding cDNA Clone identified in the second column. In the fifth column, the nucleotide sequence identified as "NT SEQ ID NO:X" was assembled from partially homologous ("overlapping") sequences obtained from the corresponding cDNA clone identified in the second column and, in some cases, from additional related cDNA clones. The overlapping sequences were assembled into a single contiguous sequence of high redundancy (usually three to five overlapping sequences at each nucleotide position), resulting in a final sequence identified as SEQ ID NO:X. In the sixth column, "Total NT Seq." refers to the total number of nucleotides in the contig sequence identified as SEQ ID NO:X." The deposited clone may contain all or most of these sequences, reflected by the nucleotide position indicated as "5" NT of Clone Seq." (seventh column) and the "3" NT of Clone Seq." (eighth column) of SEQ ID NO:X. In the ninth column, the nucleotide position of SEQ ID NO:X of the putative start codon (methionine) is identified as "5" NT of Start Codon." Similarly, in column ten, the nucleotide position of SEQ ID NO:X of the predicted signal sequence is identified as "5' NT of First AA of Signal Pep." In the eleventh column, the translated amino acid sequence, beginning with the methionine, is identified as "AA SEQ ID NO:Y," although other reading frames can also be routinely translated using known molecular biology techniques. The polypeptides produced by these alternative open reading frames are specifically contemplated by the present invention.

[9] In the twelfth and thirteenth columns of Table 1A, the first and last amino acid position of SEQ ID NO:Y of the predicted signal peptide is identified as "First AA of Sig Pep" and "Last AA of Sig Pep." In the fourteenth column, the predicted first amino acid position of SEQ ID NO:Y of the secreted portion is identified as "Predicted First AA of Secreted Portion". The amino acid position of SEQ ID NO:Y of the last amino acid encoded by the open reading frame is identified in the fifteenth column as "Last AA of ORF".

- SEQ ID NO:X (where X may be any of the polynucleotide sequences disclosed in the sequence listing) and the translated SEQ ID NO:Y (where Y may be any of the polypeptide sequences disclosed in the sequence listing) are sufficiently accurate and otherwise suitable for a variety of uses well known in the art and described further below. For instance, SEQ ID NO:X is useful for designing nucleic acid hybridization probes that will detect nucleic acid sequences contained in SEQ ID NO:X or the cDNA contained in the deposited clone. These probes will also hybridize to nucleic acid molecules in biological samples, thereby enabling a variety of forensic and diagnostic methods of the invention. Similarly, polypeptides identified from SEQ ID NO:Y may be used, for example, to generate antibodies which bind specifically to proteins containing the polypeptides and the secreted proteins encoded by the cDNA clones identified in Table 1A and/or elsewhere herein
- Nevertheless, DNA sequences generated by sequencing reactions can contain sequencing errors. The errors exist as misidentified nucleotides, or as insertions or deletions of nucleotides in the generated DNA sequence. The erroneously inserted or deleted nucleotides cause frame shifts in the reading frames of the predicted amino acid sequence. In these cases, the predicted amino acid sequence diverges from the actual amino acid sequence, even though the generated DNA sequence may be greater than 99.9% identical to the actual DNA sequence (for example, one base insertion or deletion in an open reading frame of over 1000 bases).
- [12] Accordingly, for those applications requiring precision in the nucleotide sequence or the amino acid sequence, the present invention provides not only the generated nucleotide sequence identified as SEQ ID NO:X, and the predicted translated amino acid sequence identified as SEQ ID NO:Y, but also a sample of plasmid DNA containing a human cDNA of the invention deposited with the ATCC, as set forth in Table 1A. The nucleotide sequence of each deposited plasmid can readily be determined by sequencing the deposited plasmid in accordance with known methods
- [13] The predicted amino acid sequence can then be verified from such deposits. Moreover, the amino acid sequence of the protein encoded by a particular plasmid can also be directly determined by peptide sequencing or by expressing the protein in a suitable host cell containing the deposited human cDNA, collecting the protein, and determining its sequence.

- [14] Also provided in Table 1A is the name of the vector which contains the cDNA plasmid. Each vector is routinely used in the art. The following additional information is provided for convenience.
- [15] Vectors Lambda Zap (U.S. Patent Nos. 5,128,256 and 5,286,636), Uni-Zap XR (U.S. Patent Nos. 5,128, 256 and 5,286,636), Zap Express (U.S. Patent Nos. 5,128,256 and 5,286,636), pBluescript (pBS) (Short, J. M. et al., *Nucleic Acids Res.* 16:7583-7600 (1988); Alting-Mees, M. A. and Short, J. M., *Nucleic Acids Res.* 17:9494 (1989)) and pBK (Alting-Mees, M. A. et al., *Strategies* 5:58-61 (1992)) are commercially available from Stratagene Cloning Systems, Inc., 11011 N. Torrey Pines Road, La Jolla, CA, 92037. pBS contains an ampicillin resistance gene and pBK contains a neomycin resistance gene. Phagemid pBS may be excised from the Lambda Zap and Uni-Zap XR vectors, and phagemid pBK may be excised from the Zap Express vector. Both phagemids may be transformed into *E. coli* strain XL-1 Blue, also available from Stratagene
- Vectors pSport1, pCMVSport 1.0, pCMVSport 2.0 and pCMVSport 3.0, were obtained from Life Technologies, Inc., P. O. Box 6009, Gaithersburg, MD 20897. All Sport vectors contain an ampicillin resistance gene and may be transformed into *E. coli* strain DH10B, also available from Life Technologies. See, for instance, Gruber, C. E., et al., *Focus* 15:59 (1993). Vector lafmid BA (Bento Soares, Columbia University, New York, NY) contains an ampicillin resistance gene and can be transformed into *E. coli* strain XL-1 Blue. Vector pCR<sup>®</sup>2.1, which is available from Invitrogen, 1600 Faraday Avenue, Carlsbad, CA 92008, contains an ampicillin resistance gene and may be transformed into *E. coli* strain DH10B, available from Life Technologies. See, for instance, Clark, J. M., *Nuc. Acids Res.* 16:9677-9686 (1988) and Mead, D. *et al.*, *Bio/Technology* 9: (1991).
- [17] The present invention also relates to the genes corresponding to SEQ ID NO:X, SEQ ID NO:Y, and/or a deposited cDNA (cDNA Clone ID). The corresponding gene can be isolated in accordance with known methods using the sequence information disclosed herein. Such methods include, but are not limited to, preparing probes or primers from the disclosed sequence and identifying or amplifying the corresponding gene from appropriate sources of genomic material.
- [18] Also provided in the present invention are allelic variants, orthologs, and/or species homologs. Procedures known in the art can be used to obtain full-length genes, allelic variants, splice variants, full-length coding portions, orthologs, and/or species homologs of genes corresponding to SEQ ID NO:X and SEQ ID NO:Y using information

from the sequences disclosed herein or the clones deposited with the ATCC. For example, allelic variants and/or species homologs may be isolated and identified by making suitable probes or primers from the sequences provided herein and screening a suitable nucleic acid source for allelic variants and/or the desired homologue.

The present invention provides a polynucleotide comprising, or alternatively consisting of, the nucleic acid sequence of SEQ ID NO:X and/or a cDNA contained in ATCC Deposit No.Z. The present invention also provides a polypeptide comprising, or alternatively, consisting of, the polypeptide sequence of SEQ ID NO:Y, a polypeptide encoded by SEQ ID NO:X, and/or a polypeptide encoded by a cDNA contained in ATCC deposit No.Z. Polynucleotides encoding a polypeptide comprising, or alternatively consisting of the polypeptide sequence of SEQ ID NO:Y, a polypeptide encoded by SEQ ID NO:X and/or a polypeptide encoded by the cDNA contained in ATCC Deposit No.Z, are also encompassed by the invention. The present invention further encompasses a polynucleotide comprising, or alternatively consisting of the complement of the nucleic acid sequence of SEQ ID NO:X, and/or the complement of the coding strand of the cDNA contained in ATCC Deposit No.Z.

#### **Description of Table 1B**

Table 1B summarizes some of the polynucleotides encompassed by the [20] invention (including cDNA clones related to the sequences (Clone ID NO:Z), contig sequences (contig identifier (Contig ID:) and contig nucleotide sequence identifier (SEQ ID NO:X)) and further summarizes certain characteristics of these polynucleotides and the The first column provides the gene number in the polypeptides encoded thereby. application for each clone identifier. The second column provides a unique clone identifier, "Clone ID NO:Z", for a cDNA clone related to each contig sequence disclosed in Table 1A and/or 1B. The third column provides a unique contig identifier, "Contig ID:" for each of the contig sequences disclosed in Table 1B. The fourth column provides the sequence identifier, "SEQ ID NO:X", for each of the contig sequences disclosed in Table 1A and/or 1B. The fifth column, "ORF (From-To)", provides the location (i.e., nucleotide position numbers) within the polynucleotide sequence of SEQ ID NO:X that delineate the preferred open reading frame (ORF) that encodes the amino acid sequence shown in the sequence listing and referenced in Table 1B as SEQ ID NO:Y (column 6). Column 7 lists residues comprising predicted epitopes contained in the polypeptides encoded by each of the preferred ORFs (SEQ ID NO:Y). Identification of potential immunogenic regions was performed according to the method of Jameson and Wolf (CABIOS, 4; 181-186 (1988)); specifically, the Genetics Computer Group (GCG) implementation of this algorithm, embodied in the program PEPTIDESTRUCTURE (Wisconsin Package v10.0, Genetics Computer Group (GCG), Madison, Wisc.). This method returns a measure of the probability that a given residue is found on the surface of the protein. Regions where the antigenic index score is greater than 0.9 over at least 6 amino acids are indicated in Table 1B as "Predicted Epitopes". In particular embodiments, polypeptides of the invention comprise, or alternatively consist of, one, two, three, four, five or more of the predicted epitopes described in Table 1B. It will be appreciated that depending on the analytical criteria used to predict antigenic determinants, the exact address of the determinant may vary slightly. Column 8, "Tissue Distribution" shows the expression profile of tissue, cells, and/or cell line libraries which express the polynucleotides of the invention. The first number in column 8 (preceding the colon), represents the tissue/cell source identifier code corresponding to the key provided in Table 4. Expression of these polynucleotides was not observed in the other tissues and/or cell libraries tested. For those identifier codes in which the first two letters are not "AR", the second number in column 8 (following the colon), represents the number of times a sequence corresponding to the reference polynucleotide sequence (e.g., SEQ ID NO:X) was identified in the tissue/cell source. Those tissue/cell source identifier codes in which the first two letters are "AR" designate information generated using DNA array Utilizing this technology, cDNAs were amplified by PCR and then technology. transferred, in duplicate, onto the array. Gene expression was assayed through hybridization of first strand cDNA probes to the DNA array. cDNA probes were generated from total RNA extracted from a variety of different tissues and cell lines. Probe synthesis was performed in the presence of <sup>33</sup>P dCTP, using oligo(dT) to prime reverse transcription. After hybridization, high stringency washing conditions were employed to remove non-specific hybrids from the array. The remaining signal, emanating from each gene target, was measured using a Phosphorimager. Gene expression was reported as Phosphor Stimulating Luminescence (PSL) which reflects the level of phosphor signal generated from the probe hybridized to each of the gene targets represented on the array. A local background signal subtraction was performed before the total signal generated from each array was used to normalize gene expression between the different hybridizations. The value presented after "[array code]:" represents the mean of the duplicate values, following background subtraction and probe normalization. One of skill

in the art could routinely use this information to identify normal and/or diseased tissue(s) which show a predominant expression pattern of the corresponding polynucleotide of the invention or to identify polynucleotides which show predominant and/or specific tissue and/or cell expression. Column 9 provides the chromosomal location of polynucleotides corresponding to SEQ ID NO:X. Chromosomal location was determined by finding exact matches to EST and cDNA sequences contained in the NCBI (National Center for Biotechnology Information) UniGene database. Given a presumptive chromosomal location, disease locus association was determined by comparison with the Morbid Map, derived from Online Mendelian Inheritance in Man (Online Mendelian Inheritance in Man, OMIM<sup>TM</sup>. McKusick-Nathans Institute for Genetic Medicine, Johns Hopkins University (Baltimore, MD) and National Center for Biotechnology Information, National MD) 2000. World Wide Web URL: (Bethesda, Library of Medicine http://www.ncbi.nlm.nih.gov/omim/). If the putative chromosomal location of the Query overlaps with the chromosomal location of a Morbid Map entry, an OMIM identification number is disclosed in column 10 labeled "OMIM Disease Reference(s)". A key to the OMIM reference identification numbers is provided in Table 5.

### **Description of Table 1C**

Table 1C summarizes additional polynucleotides encompassed by the invention [21] (including cDNA clones related to the sequences (Clone ID NO:Z), contig sequences (contig identifier (Contig ID:) contig nucleotide sequence identifiers (SEQ ID NO:X)), and genomic sequences (SEQ ID NO:B). The first column provides a unique clone identifier, "Clone ID NO:Z", for a cDNA clone related to each contig sequence. The second column provides the sequence identifier, "SEQ ID NO:X", for each contig sequence. The third column provides a unique contig identifier, "Contig ID:" for each contig sequence. The fourth column, provides a BAC identifier "BAC ID NO:A" for the BAC clone referenced in the corresponding row of the table. The fifth column provides the nucleotide sequence identifier, "SEQ ID NO:B" for a fragment of the BAC clone identified in column four of the corresponding row of the table. The sixth column, "Exon From-To", provides the location (i.e., nucleotide position numbers) within the polynucleotide sequence of SEQ ID NO:B which delineate certain polynucleotides of the invention that are also exemplary members of polynucleotide sequences that encode polypeptides of the invention (e.g., polypeptides containing amino acid sequences encoded by the polynucleotide sequences delineated in column six, and fragments and variants thereof).

## **Description of Table 1D**

Table 1D: In preferred embodiments, the present invention encompasses a method of treating a disease or disorder listed in the "FEATURES OF PROTEIN" sections (below) and also as listed in the "Preferred Indications" column of Table 1D (below); comprising administering to a patient in which such treatment, prevention, or amelioration is desired a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) represented by Table 1A and Table 1D (in the same row as the disease or disorder to be treated is listed in the "Preferred Indications" column of Table 1D) in an amount effective to treat, prevent, or ameliorate the disease or disorder.

[23] As indicated in Table 1D, the polynucleotides, polypeptides, agonists, or antagonists of the present invention (including antibodies) can be used in assays to test for one or more biological activities. If these polynucleotides and polypeptides do exhibit activity in a particular assay, it is likely that these molecules may be involved in the diseases associated with the biological activity. Thus, the polynucleotides or polypeptides, or agonists or antagonists thereof (including antibodies) could be used to treat the associated disease.

The present invention encompasses methods of preventing, treating, diagnosing, or ameliorating a disease or disorder. In preferred embodiments, the present invention encompasses a method of treating a disease or disorder listed in the "Preferred Indications" column of Table 1D; comprising administering to a patient in which such treatment, prevention, or amelioration is desired a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) in an amount effective to treat, prevent, diagnose, or ameliorate the disease or disorder. The first and seccond columns of Table 1D show the "Gene No." and "cDNA Clone ID No.", respectively, indicating certain nucleic acids and proteins (or antibodies against the same) of the invention (including polynucleotide, polypeptide, and antibody fragments or variants thereof) that may be used in preventing, treating, diagnosing, or ameliorating the disease(s) or disorder(s) indicated in the corresponding row in Column 3 of Table 1D.

[25] In another embodiment, the present invention also encompasses methods of preventing, treating, diagnosing, or ameliorating a disease or disorder listed in the "Preferred Indications" column of Table 1D; comprising administering to a patient

combinations of the proteins, nucleic acids, or antibodies of the invention (or fragments or variants thereof), sharing similar indications as shown in the corresponding rows in Column 3 of Table 1D.

[26] The "Preferred Indication" column describes diseases, disorders, and/or conditions that may be treated, prevented, diagnosed, or ameliorated by a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof).

The recitation of "Cancer" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof) may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., leukemias, cancers, and/or as described below under "Hyperproliferative Disorders").

In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Cancer" recitation in the "Preferred Indication" column of Table 1D may be used for example, to diagnose, treat, prevent, and/or ameliorate a neoplasm located in a tissue selected from the group consisting of: colon, abdomen, bone, breast, digestive system, liver, pancreas, prostate, peritoneum, lung, blood (e.g., leukemia), endocrine glands (adrenal, parathyroid, pituitary, testicles, ovary, thymus, thyroid), uterus, eye, head and neck, nervous (central and peripheral), lymphatic system, pelvic, skin, soft tissue, spleen, thoracic, and urogenital.

In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Cancer" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a pre-neoplastic condition, selected from the group consisting of: hyperplasia (e.g., endometrial hyperplasia and/or as described in the section entitled "Hyperproliferative Disorders"), metaplasia (e.g., connective tissue metaplasia, atypical metaplasia, and/or as described in the section entitled "Hyperproliferative Disorders"), and/or dysplasia (e.g., cervical dysplasia, and bronchopulmonary dysplasia).

[30] In another specific embodiment, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Cancer" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a benign dysproliferative disorder selected from the group consisting of: benign tumors, fibrocystic conditions, tissue hypertrophy, and/or as described in the section entitled "Hyperproliferative Disorders".

- [31] The recitation of "Immune/Hematopoietic" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders"), blood disorders (e.g., as described below under "Immune Activity" "Cardiovascular Disorders" and/or "Blood-Related Disorders"), and infections (e.g., as described below under "Infectious Disease").
- In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having the "Immune/Hematopoietic" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: anemia, pancytopenia, leukopenia, thrombocytopenia, leukemias, Hodgkin's disease, non-Hodgkin's lymphoma, acute lymphocytic anemia (ALL), plasmacytomas, multiple myeloma, Burkitt's lymphoma, arthritis, asthma, AIDS, autoimmune disease, rheumatoid arthritis, granulomatous disease, immune deficiency, inflammatory bowel disease, sepsis, neutropenia, neutrophilia, psoriasis, immune reactions to transplanted organs and tissues, systemic lupus erythematosis, hemophilia, hypercoagulation, diabetes mellitus, endocarditis, meningitis, Lyme Disease, and allergies.
- [33] The recitation of "Reproductive" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders"), and disorders of the reproductive system (e.g., as described below under "Reproductive System Disorders").
- [34] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Reproductive" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: cryptorchism, prostatitis, inguinal hernia, varicocele, leydig cell tumors, verrucous carcinoma, prostatitis, malacoplakia, Peyronie's disease, penile carcinoma, squamous cell hyperplasia, dysmenorrhea, ovarian adenocarcinoma, Turner's syndrome, mucopurulent cervicitis, Sertoli-leydig tumors, ovarian cancer, uterine cancer, pelvic inflammatory disease, testicular cancer, prostate cancer, Klinefelter's syndrome, Young's syndrome,

premature ejaculation, diabetes mellitus, cystic fibrosis, Kartagener's syndrome, testicular atrophy, testicular feminization, anorchia, ectopic testis, epididymitis, orchitis, gonorrhea, syphilis, testicular torsion, vasitis nodosa, germ cell tumors, stromal tumors, dysmenorrhea, retroverted uterus, endometriosis, fibroids, adenomyosis, anovulatory bleeding, amenorrhea, Cushing's syndrome, hydatidiform moles, Asherman's syndrome, premature menopause, precocious puberty, uterine polyps, dysfunctional uterine bleeding, cervicitis, chronic cervicitis, mucopurulent cervicitis, cervical dysplasia, cervical polyps, Nabothian cysts, cervical erosion, cervical incompetence, cervical neoplasms, pseudohermaphroditism, and premenstrual syndrome.

- [35] The recitation of "Musculoskeletal" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders"), and disorders of the immune system (e.g., as described below under "Immune Activity").
- [36] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Musculoskeletal" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: bone cancers (e.g., osteochondromas, benign chondromas, chondroblastoma, chondromyxoid fibromas, osteoid osteomas, giant cell tumors, multiple myeloma, osteosarcomas), Paget's Disease, rheumatoid arthritis, systemic lupus erythematosus, osteomyelitis, Lyme Disease, gout, bursitis, tendonitis, osteoporosis, osteoarthritis, muscular dystrophy, mitochondrial myopathy, cachexia, and multiple sclerosis.
- [37] The recitation of "Cardiovascular" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders"), and disorders of the cardiovascular system (e.g., as described below under "Cardiovascular Disorders").
- [38] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Cardiovascular" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: myxomas,

fibromas, rhabdomyomas, cardiovascular abnormalities (e.g., congenital heart defects, cerebral arteriovenous malformations, septal defects), heart disease (e.g., heart failure, congestive heart disease, arrhythmia, tachycardia, fibrillation, pericardial Disease, endocarditis), cardiac arrest, heart valve disease (e.g., stenosis, regurgitation, prolapse), vascular disease (e.g., hypertension, coronary artery disease, angina, aneurysm, arteriosclerosis, peripheral vascular disease), hyponatremia, hypernatremia, hypokalemia, and hyperkalemia.

- [39] The recitation of "Mixed Fetal" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders").
- In specific embodiments, a protein, nucleic acid, or antibody of the invention [40] (or fragment or variant thereof) having a "Mixed Fetal" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: spina bifida, hydranencephaly, neurofibromatosis, fetal alcohol syndrome, diabetes mellitus, PKU, Down's syndrome, Patau syndrome, Edwards syndrome, Turner syndrome, Apert syndrome, Carpenter syndrome, Conradi syndrome, Crouzon syndrome, cutis laxa, Cornelia de Lange syndrome, Ellis-van Creveld syndrome, Holt-Oram syndrome, Kartagener syndrome, Meckel-Gruber syndrome, Noonan syndrome, Pallister-Hall syndrome, Rubinstein-Taybi syndrome, Scimitar syndrome, Smith-Lemli-Opitz syndrome, thromocytopenia-absent radius (TAR) syndrome, Treacher Collins syndrome, Williams syndrome, Hirschsprung's disease, Meckel's diverticulum, polycystic kidney disease, Turner's syndrome, and gonadal dysgenesis, Klippel-Feil syndrome, Ostogenesis imperfecta, muscular dystrophy, Tay-Sachs disease, Wilm's tumor, neuroblastoma, and retinoblastoma.
- [41] The recitation of "Excretory" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders") and renal disorders (e.g., as described below under "Renal Disorders").

- In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Excretory" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: bladder cancer, prostate cancer, benign prostatic hyperplasia, bladder disorders (e.g., urinary incontinence, urinary retention, urinary obstruction, urinary tract Infections, interstitial cystitis, prostatitis, neurogenic bladder, hematuria), renal disorders (e.g., hydronephrosis, proteinuria, renal failure, pyelonephritis, urolithiasis, reflux nephropathy, and unilateral obstructive uropathy).
- [43] The recitation of "Neural/Sensory" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders") and diseases or disorders of the nervous system (e.g., as described below under "Neural Activity and Neurological Diseases").
- In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Neural/Sensory" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: brain cancer (e.g., brain stem glioma, brain tumors, central nervous system (Primary) lymphoma, central nervous system lymphoma, cerebellar astrocytoma, and cerebral astrocytoma, neurodegenerative disorders (e.g., Alzheimer's Disease, Creutzfeldt-Jakob Disease, Parkinson's Disease, and Idiopathic Presenile Dementia), encephalomyelitis, cerebral malaria, meningitis, metabolic brain diseases (e.g., phenylketonuria and pyruvate carboxylase deficiency), cerebellar ataxia, ataxia telangiectasia, and AIDS Dementia Complex, schizophrenia, attention deficit disorder, hyperactive attention deficit disorder, autism, and obsessive compulsive disorders.
- [45] The recitation of "Respiratory" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders") and diseases or disorders of the respiratory system (e.g., as described below under "Respiratory Disorders").

In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Respiratory" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: cancers of the respiratory system such as larynx cancer, pharynx cancer, trachea cancer, epiglottis cancer, lung cancer, squamous cell carcinomas, small cell (oat cell) carcinomas, large cell carcinomas, and adenocarcinomas. Allergic reactions, cystic fibrosis, sarcoidosis, histiocytosis X, infiltrative lung diseases (e.g., pulmonary fibrosis and lymphoid interstitial pneumonia), obstructive airway diseases (e.g., asthma, emphysema, chronic or acute bronchitis), occupational lung diseases (e.g., silicosis and asbestosis), pneumonia, and pleurisy.

The recitation of "Endocrine" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders") and diseases or disorders of the respiratory system (e.g., as described below under "Respiratory Disorders"), renal disorders (e.g., as described below under "Renal Disorders"), and disorders of the endocrine system (e.g., as described below under "Endocrine Disorders".

[48] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having an "Endocrine" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: cancers of endocrine tissues and organs (e.g., cancers of the hypothalamus, pituitary gland, thyroid gland, parathyroid glands, pancreas, adrenal glands, ovaries, and testes), diabetes (e.g., diabetes insipidus, type I and type II diabetes mellitus), obesity, disorders related to pituitary glands (e.g., hyperpituitarism, hypopituitarism, and pituitary dwarfism), hypothyroidism, hyperthyroidism, goiter, reproductive disorders (e.g. male and female infertility), disorders related to adrenal glands (e.g., Addison's Disease, corticosteroid deficiency, and Cushing's Syndrome), kidney cancer (e.g., hypernephroma, transitional cell cancer, and Wilm's tumor), diabetic nephropathy, interstitial nephritis, polycystic kidney disease, glomerulonephritis (e.g., IgM mesangial proliferative glomerulonephritis and glomerulonephritis caused by autoimmune disorders; such as Goodpasture's syndrome), and nephrocalcinosis.

- [49] The recitation of "Digestive" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders") and diseases or disorders of the gastrointestinal system (e.g., as described below under "Gastrointestinal Disorders".
- In specific embodiments, a protein, nucleic acid, or antibody of the invention [50] (or fragment or variant thereof) having a "Digestive" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: ulcerative colitis, appendicitis, Crohn's disease, hepatitis, hepatic encephalopathy, portal hypertension, cholelithiasis, cancer of the digestive system (e.g., biliary tract cancer, stomach cancer, colon cancer, gastric cancer, pancreatic cancer, cancer of the bile duct, tumors of the colon (e.g., polyps or cancers), and cirrhosis), pancreatitis, ulcerative disease, pyloric stenosis, gastroenteritis, gastritis, gastric atropy, benign tumors of the duodenum, distension, irritable bowel syndrome, malabsorption, congenital disorders of the small intestine, bacterial and parasitic infection, megacolon, Hirschsprung's disease, aganglionic megacolon, acquired megacolon, colitis, anorectal disorders (e.g., anal fistulas, hemorrhoids), congenital disorders of the liver (e.g., Wilson's disease, hemochromatosis, cystic fibrosis, biliary atresia, and alpha1-antitrypsin deficiency), portal hypertension, cholelithiasis, and jaundice.
- [51] The recitation of "Connective/Epithelial" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders"), cellular and genetic abnormalities (e.g., as described below under "Diseases at the Cellular Level "), angiogenesis (e.g., as described below under "Anti-Angiogenesis Activity "), and or to promote or inhibit regeneration (e.g., as described below under "Regeneration "), and wound healing (e.g., as described below under "Wound Healing and Epithelial Cell Proliferation").
- [52] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Connective/Epithelial" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of:

metaplasia, mixed connective tissue disease, focal epithelial hyperplasia, epithelial metaplasia, mucoepithelial dysplasia, graft v. host disease, polymyositis, cystic hyperplasia, cerebral dysplasia, tissue hypertrophy, Alzheimer's disease, lymphoproliferative disorder, Waldenstron's macroglobulinemia, Crohn's disease, pernicious anemia, idiopathic Addison's disease, glomerulonephritis, bullous pemphigoid, Sjogren's syndrome, diabetes mellitus, cystic fibrosis, osteoblastoma, osteoclastoma, osteosarcoma, chondrosarcoma, osteoporosis, osteocarthritis, periodontal disease, wound healing, relapsing polychondritis, vasculitis, polyarteritis nodosa, Wegener's granulomatosis, cellulitis, rheumatoid arthritis, psoriatic arthritis, discoid lupus erythematosus, systemic lupus erythematosus, scleroderma, CREST syndrome, Sjogren's syndrome, polymyositis, dermatomyositis, mixed connective tissue disease, relapsing polychondritis, vasculitis, Henoch-Schonlein syndrome, erythema nodosum, polyarteritis nodosa, temporal (giant cell) arteritis, Takayasu's arteritis, Wegener's granulomatosis, Reiter's syndrome, Behcet's syndrome, ankylosing spondylitis, cellulitis, keloids, Ehler Danlos syndrome, Marfan syndrome, pseudoxantoma elasticum, osteogenese imperfecta, chondrodysplasias, epidermolysis bullosa, Alport syndrome, and cutis laxa.

#### **Description of Table 1E**

Table 1E provides information related to biological activities and preferred [53] indications for polynucleotides and polypeptides of the invention (including antibodies, agonists, and/or antagonists thereof). Table 1E also provides information related to assays which may be used to test polynucleotides and polypeptides of the invention (including antibodies, agonists, and/or antagonists thereof) for the corresponding biological activities. The first column ("Gene No.") provides the gene number in the application for each clone identifier. The second column ("cDNA Clone ID No:Z") provides the unique clone identifier for each clone as previously described and indicated in Tables 1A, 1B, 1C, and 1D. The third column ("AA SEQ ID NO:Y") indicates the Sequence Listing SEQ ID Number for polypeptide sequences encoded by the corresponding cDNA clones (also as indicated in Tables 1A, 1B, and 2). The fourth column ("Biological Activity") indicates a biological activity corresponding to the indicated polypeptides (or polynucleotides encoding said The fifth column ("Exemplary Activity Assay") further describes the polypeptides). corresponding biological activity and provides information pertaining to the various types of assays which may be performed to test, demonstrate, or quantify the corresponding biological activity. The sixth column ("Preferred Indications") describes particular embodiments of the invention and indications (e.g. pathologies, diseases, disorders, abnormalities, etc.) for which polynucleotides and polypeptides of the invention (including antibodies, agonists, and/or antagonists thereof) may be used in detecting, diagnosing, preventing, and/or treating.

Table 1E describes the use of FMAT technology, inter alia, for testing or [54] demonstrating various biological activities. Fluorometric microvolume assay technology (FMAT) is a fluorescence-based system which provides a means to perform nonradioactive cell- and bead-based assays to detect activation of cell signal transduction pathways. This technology was designed specifically for ligand binding and immunological assays. Using this technology, fluorescent cells or beads at the bottom of the well are detected as localized areas of concentrated fluorescence using a data processing system. Unbound flurophore comprising the background signal is ignored, allowing for a wide variety of homogeneous assays. FMAT technology may be used for peptide ligand binding assays, immunofluorescence, apoptosis, cytotoxicity, and beadbased immunocapture assays. See, Miraglia S et. al., "Homogeneous cell and bead based assays for highthroughput screening using flourometric microvolume assay technology," Journal of Biomolecular Screening; 4:193-204 (1999). In particular, FMAT technology may be used to test, confirm, and/or identify the ability of polypeptides (including polypeptide fragments and variants) to activate signal transduction pathways. example, FMAT technology may be used to test, confirm, and/or identify the ability of polypeptides to upregulate production of immunomodulatory proteins (such as, for example, interleukins, GM-CSF, Rantes, and Tumor Necrosis factors, as well as other cellular regulators (e.g. insulin)).

Table 1E also describes the use of kinase assays for testing, demonstrating, or quantifying biological activity. In this regard, the phosphorylation and dephosphorylation of specific amino acid residues (e.g. Tyrosine, Serine, Threonine) on cell-signal transduction proteins provides a fast, reversible means for activation and deactivation of cellular signal transduction pathways. Moreover, cell signal transduction via phosphorylation/de-phosphorylation is crucial to the regulation of a wide variety of cellular processes (e.g. proliferation, differentiation, migration, apoptosis, etc.). Accordingly, kinase assays provide a powerful tool useful for testing, confirming, and/or identifying polypeptides (including polypeptide fragments and variants) that mediate cell signal transduction events via protein phosphorylation. See e.g., Forrer, P., Tamaskovic R., and Jaussi, R. "Enzyme-Linked Immunosorbent Assay for Measurement of JNK, ERK, and p38 Kinase Activities" Biol. Chem. 379(8-9): 1101-1110 (1998).

#### **Description of Table 2**

[56] Table 2 summarizes homology and features of some of the polypeptides of the invention. The first column provides a unique clone identifier, "Clone ID NO:Z", corresponding to a cDNA clone disclosed in Table 1A or 1B. The second column provides the unique contig identifier, "Contig ID:" corresponding to contigs in Table 1B and allowing for correlation with the information in Table 1B. The third column provides the sequence identifier, "SEQ ID NO:X", for the contig polynucleotide sequence. The fourth column provides the analysis method by which the homology/identity disclosed in the Table was determined. Comparisons were made between polypeptides encoded by the polynucleotides of the invention and either a non-redundant protein database (herein referred to as "NR"), or a database of protein families (herein referred to as "PFAM") as further described below. The fifth column provides a description of the PFAM/NR hit having a significant match to a polypeptide of the invention. Column six provides the accession number of the PFAM/NR hit disclosed in the fifth column. Column seven, "Score/Percent Identity", provides a quality score or the percent identity, of the hit disclosed in columns five and six. Columns 8 and 9, "NT From" and "NT To" respectively, delineate the polynucleotides in "SEQ ID NO:X" that encode a polypeptide having a significant match to the PFAM/NR database as disclosed in the fifth and sixth In specific embodiments polypeptides of the invention comprise, or columns. alternatively consist of, an amino acid sequence encoded by a polynucleotide in SEQ ID NO:X as delineated in columns 8 and 9, or fragments or variants thereof.

#### **Description of Table 3**

Table 3 provides polynucleotide sequences that may be disclaimed according to certain embodiments of the invention. The first column provides a unique clone identifier, "Clone ID", for a cDNA clone related to contig sequences disclosed in Table 1B. The second column provides the sequence identifier, "SEQ ID NO:X", for contig sequences disclosed in Table 1A and/or 1B. The third column provides the unique contig identifier,

"Contig ID:", for contigs disclosed in Table 1B. The fourth column provides a unique integer 'a' where 'a' is any integer between 1 and the final nucleotide minus 15 of SEQ ID NO:X, and the fifth column provides a unique integer 'b' where 'b' is any integer between 15 and the final nucleotide of SEQ ID NO:X, where both a and b correspond to the positions of nucleotide residues shown in SEQ ID NO:X, and where b is greater than or equal to a + 14. For each of the polynucleotides shown as SEQ ID NO:X, the uniquely defined integers can be substituted into the general formula of a-b, and used to describe polynucleotides which may be preferably excluded from the invention. In certain embodiments, preferably excluded from the invention are at least one, two, three, four, five, ten, or more of the polynucleotide sequence(s) having the accession number(s) disclosed in the sixth column of this Table (including for example, published sequence in connection with a particular BAC clone). In further embodiments, preferably excluded from the invention are the specific polynucleotide sequence(s) contained in the clones corresponding to at least one, two, three, four, five, ten, or more of the available material having the accession numbers identified in the sixth column of this Table (including for example, the actual sequence contained in an identified BAC clone).

## **Description of Table 4**

Table 4 provides a key to the tissue/cell source identifier code disclosed in Table 1B, column 8. Column 1 provides the tissue/cell source identifier code disclosed in Table 1B, Column 8. Columns 2-5 provide a description of the tissue or cell source. Codes corresponding to diseased tissues are indicated in column 6 with the word "disease". The use of the word "disease" in column 6 is non-limiting. The tissue or cell source may be specific (e.g. a neoplasm), or may be disease-associated (e.g., a tissue sample from a normal portion of a diseased organ). Furthermore, tissues and/or cells lacking the "disease" designation may still be derived from sources directly or indirectly involved in a disease state or disorder, and therefore may have a further utility in that disease state or disorder. In numerous cases where the tissue/cell source is a library, column 7 identifies the vector used to generate the library.

#### **Description of Table 5**

[59] Table 5 provides a key to the OMIM reference identification numbers disclosed in Table 1B, column 10. OMIM reference identification numbers (Column 1) were derived from Online Mendelian Inheritance in Man (Online Mendelian Inheritance in

Man, OMIM. McKusick-Nathans Institute for Genetic Medicine, Johns Hopkins University (Baltimore, MD) and National Center for Biotechnology Information, National Library of Medicine, (Bethesda, MD) 2000. World Wide Web URL: http://www.ncbi.nlm.nih.gov/omim/). Column 2 provides diseases associated with the cytologic band disclosed in Table 1B, column 9, as determined using the Morbid Map database.

#### **Description of Table 6**

[60] Table 6 summarizes some of the ATCC Deposits, Deposit dates, and ATCC designation numbers of deposits made with the ATCC in connection with the present application. These deposits were made in addition to those described in the Table 1A.

### **Description of Table 7**

- [61] Table 7 shows the cDNA libraries sequenced, and ATCC designation numbers and vector information relating to these cDNA libraries.
- [62] The first column shows the first four letters indicating the Library from which each library clone was derived. The second column indicates the catalogued tissue description for the corresponding libraries. The third column indicates the vector containing the corresponding clones. The fourth column shows the ATCC deposit designation for each library clone as indicated by the deposit information in Table 6.

## **Definitions**

- [63] The following definitions are provided to facilitate understanding of certain terms used throughout this specification.
- In the present invention, "isolated" refers to material removed from its original environment (e.g., the natural environment if it is naturally occurring), and thus is altered "by the hand of man" from its natural state. For example, an isolated polynucleotide could be part of a vector or a composition of matter, or could be contained within a cell, and still be "isolated" because that vector, composition of matter, or particular cell is not the original environment of the polynucleotide. The term "isolated" does not refer to genomic or cDNA libraries, whole cell total or mRNA preparations, genomic DNA preparations

(including those separated by electrophoresis and transferred onto blots), sheared whole cell genomic DNA preparations or other compositions where the art demonstrates no distinguishing features of the polynucleotide/sequences of the present invention.

- In the present invention, a "secreted" protein refers to those proteins capable of being directed to the ER, secretory vesicles, or the extracellular space as a result of a signal sequence, as well as those proteins released into the extracellular space without necessarily containing a signal sequence. If the secreted protein is released into the extracellular space, the secreted protein can undergo extracellular processing to produce a "mature" protein. Release into the extracellular space can occur by many mechanisms, including exocytosis and proteolytic cleavage.
- As used herein, a "polynucleotide" refers to a molecule having a nucleic acid [66] sequence encoding SEQ ID NO:Y or a fragment or variant thereof (e.g., the polypeptide delinated in columns fourteen and fifteen of Table 1A); a nucleic acid sequence contained in SEQ ID NO:X (as described in column 5 of Table 1A and/or column 3 of Table 1B) or the complement thereof; a cDNA sequence contained in Clone ID NO:Z (as described in column 2 of Table 1A and/or 1B and contained within a library deposited with the ATCC); a nucleotide sequence encoding the polypeptide encoded by a nucleotide sequence in SEQ ID NO:B as defined in column 6 (EXON From-To) of Table 1C or a fragment or variant thereof; or a nucleotide coding sequence in SEQ ID NO:B as defined in column 6 of Table 1C or the complement thereof. For example, the polynucleotide can contain the nucleotide sequence of the full length cDNA sequence, including the 5' and 3' untranslated sequences, the coding region, as well as fragments, epitopes, domains, and variants of the nucleic acid sequence. Moreover, as used herein, a "polypeptide" refers to a molecule having an amino acid sequence encoded by a polynucleotide of the invention as broadly defined (obviously excluding poly-Phenylalanine or poly-Lysine peptide sequences which result from translation of a polyA tail of a sequence corresponding to a cDNA).
- [67] In the present invention, "SEQ ID NO:X" was often generated by overlapping sequences contained in multiple clones (contig analysis). A representative clone containing all or most of the sequence for SEQ ID NO:X is deposited at Human Genome Sciences, Inc. (HGS) in a catalogued and archived library. As shown, for example, in

column 2 of Table 1B, each clone is identified by a cDNA Clone ID (identifier generally referred to herein as Clone ID NO:Z). Each Clone ID is unique to an individual clone and the Clone ID is all the information needed to retrieve a given clone from the HGS library. Table 7 provides a list of the deposited cDNA libraries. One can use the Clone ID NO:Z to determine the library source by reference to Tables 6 and 7. Table 7 lists the deposited cDNA libraries by name and links each library to an ATCC Deposit. Library names contain four characters, for example, "HTWE." The name of a cDNA clone (Clone ID) isolated from that library begins with the same four characters, for example "HTWEP07". As mentioned below, Table 1A and/or 1B correlates the Clone ID names with SEQ ID NO:X. Thus, starting with an SEQ ID NO:X, one can use Tables 1A, 1B, 6, 7, and 9 to determine the corresponding Clone ID, which library it came from and which ATCC deposit the library is contained in. Furthermore, it is possible to retrieve a given cDNA clone from the source library by techniques known in the art and described elsewhere herein. The ATCC is located at 10801 University Boulevard, Manassas, Virginia 20110-2209, USA. The ATCC deposits were made pursuant to the terms of the Budapest Treaty on the international recognition of the deposit of microorganisms for the purposes of patent procedure.

[68] In specific embodiments, the polynucleotides of the invention are at least 15, at least 30, at least 50, at least 100, at least 125, at least 500, or at least 1000 continuous nucleotides but are less than or equal to 300 kb, 200 kb, 100 kb, 50 kb, 15 kb, 10 kb, 7.5kb, 5 kb, 2.5 kb, 2.0 kb, or 1 kb, in length. In a further embodiment, polynucleotides of the invention comprise a portion of the coding sequences, as disclosed herein, but do not comprise all or a portion of any intron. In another embodiment, the polynucleotides comprising coding sequences do not contain coding sequences of a genomic flanking gene (i.e., 5' or 3' to the gene of interest in the genome). In other embodiments, the polynucleotides of the invention do not contain the coding sequence of more than 1000, 500, 250, 100, 50, 25, 20, 15, 10, 5, 4, 3, 2, or 1 genomic flanking gene(s).

[69] A "polynucleotide" of the present invention also includes those polynucleotides capable of hybridizing, under stringent hybridization conditions, to sequences contained in SEQ ID NO:X, or the complement thereof (e.g., the complement of any one, two, three, four, or more of the polynucleotide fragments described herein), the polynucleotide sequence delineated in columns 7 and 8 of Table 1A or the complement thereof, the

polynucleotide sequence delineated in columns 8 and 9 of Table 2 or the complement thereof, and/or cDNA sequences contained in Clone ID NO:Z (e.g., the complement of any one, two, three, four, or more of the polynucleotide fragments, or the cDNA clone within the pool of cDNA clones deposited with the ATCC, described herein), and/or the polynucleotide sequence delineated in column 6 of Table 1C or the complement thereof. "Stringent hybridization conditions" refers to an overnight incubation at 42 degree C in a solution comprising 50% formamide, 5x SSC (750 mM NaCl, 75 mM trisodium citrate), 50 mM sodium phosphate (pH 7.6), 5x Denhardt's solution, 10% dextran sulfate, and 20  $\mu$ g/ml denatured, sheared salmon sperm DNA, followed by washing the filters in 0.1x SSC at about 65 degree C.

- [70] Also contemplated are nucleic acid molecules that hybridize to the polynucleotides of the present invention at lower stringency hybridization conditions. Changes in the stringency of hybridization and signal detection are primarily accomplished through the manipulation of formamide concentration (lower percentages of formamide result in lowered stringency); salt conditions, or temperature. For example, lower stringency conditions include an overnight incubation at 37 degree C in a solution comprising 6X SSPE (20X SSPE = 3M NaCl; 0.2M NaH<sub>2</sub>PO<sub>4</sub>; 0.02M EDTA, pH 7.4), 0.5% SDS, 30% formamide, 100 ug/ml salmon sperm blocking DNA; followed by washes at 50 degree C with 1XSSPE, 0.1% SDS. In addition, to achieve even lower stringency, washes performed following stringent hybridization can be done at higher salt concentrations (e.g. 5X SSC).
- [71] Note that variations in the above conditions may be accomplished through the inclusion and/or substitution of alternate blocking reagents used to suppress background in hybridization experiments. Typical blocking reagents include Denhardt's reagent, BLOTTO, heparin, denatured salmon sperm DNA, and commercially available proprietary formulations. The inclusion of specific blocking reagents may require modification of the hybridization conditions described above, due to problems with compatibility.
- [72] Of course, a polynucleotide which hybridizes only to polyA+ sequences (such as any 3' terminal polyA+ tract of a cDNA shown in the sequence listing), or to a complementary stretch of T (or U) residues, would not be included in the definition of

"polynucleotide," since such a polynucleotide would hybridize to any nucleic acid molecule containing a poly (A) stretch or the complement thereof (e.g., practically any double-stranded cDNA clone generated using oligo dT as a primer).

The polynucleotide of the present invention can be composed of any polyribonucleotide or polydeoxribonucleotide, which may be unmodified RNA or DNA or modified RNA or DNA. For example, polynucleotides can be composed of single- and double-stranded DNA, DNA that is a mixture of single- and double-stranded regions, single- and double-stranded RNA, and RNA that is mixture of single- and double-stranded regions, hybrid molecules comprising DNA and RNA that may be single-stranded or, more typically, double-stranded or a mixture of single- and double-stranded regions. In addition, the polynucleotide can be composed of triple-stranded regions comprising RNA or DNA or both RNA and DNA. A polynucleotide may also contain one or more modified bases or DNA or RNA backbones modified for stability or for other reasons. "Modified" bases include, for example, tritylated bases and unusual bases such as inosine. A variety of modifications can be made to DNA and RNA; thus, "polynucleotide" embraces chemically, enzymatically, or metabolically modified forms.

In specific embodiments, the polynucleotides of the invention are at least 15, at least 30, at least 50, at least 100, at least 125, at least 500, or at least 1000 continuous nucleotides but are less than or equal to 300 kb, 200 kb, 100 kb, 50 kb, 15 kb, 10 kb, 7.5kb, 5 kb, 2.5 kb, 2.0 kb, or 1 kb, in length. In a further embodiment, polynucleotides of the invention comprise a portion of the coding sequences, as disclosed herein, but do not comprise all or a portion of any intron. In another embodiment, the polynucleotides comprising coding sequences do not contain coding sequences of a genomic flanking gene (i.e., 5' or 3' to the gene of interest in the genome). In other embodiments, the polynucleotides of the invention do not contain the coding sequence of more than 1000, 500, 250, 100, 50, 25, 20, 15, 10, 5, 4, 3, 2, or 1 genomic flanking gene(s).

"SEQ ID NO:X" refers to a polynucleotide sequence described in column 5 of Table 1A, while "SEQ ID NO:Y" refers to a polypeptide sequence described in column 10 of Table 1A. SEQ ID NO:X is identified by an integer specified in column 6 of Table 1A. The polypeptide sequence SEQ ID NO:Y is a translated open reading frame (ORF) encoded by polynucleotide SEQ ID NO:X. The polynucleotide sequences are shown in the

sequence listing immediately followed by all of the polypeptide sequences. Thus, a polypeptide sequence corresponding to polynucleotide sequence SEQ ID NO:2 is the first polypeptide sequence shown in the sequence listing. The second polypeptide sequence corresponds to the polynucleotide sequence shown as SEQ ID NO:3, and so on.

**[76]** The polypeptide of the present invention can be composed of amino acids joined to each other by peptide bonds or modified peptide bonds, i.e., peptide isosteres, and may contain amino acids other than the 20 gene-encoded amino acids. polypeptides may be modified by either natural processes, such as posttranslational processing, or by chemical modification techniques which are well known in the art. Such modifications are well described in basic texts and in more detailed monographs, as well Modifications can occur anywhere in a as in a voluminous research literature. polypeptide, including the peptide backbone, the amino acid side-chains and the amino or carboxyl termini. It will be appreciated that the same type of modification may be present in the same or varying degrees at several sites in a given polypeptide. Also, a given polypeptide may contain many types of modifications. Polypeptides may be branched, for example, as a result of ubiquitination, and they may be cyclic, with or without branching. Cyclic, branched, and branched cyclic polypeptides may result from posttranslation natural processes or may be made by synthetic methods. Modifications include acetylation, acylation, ADP-ribosylation, amidation, covalent attachment of flavin, covalent attachment of a heme moiety, covalent attachment of a nucleotide or nucleotide derivative, covalent attachment of a lipid or lipid derivative, covalent attachment of phosphotidylinositol, cross-linking, cyclization, disulfide bond formation, demethylation, formation of covalent cross-links, formation of cysteine, formation of pyroglutamate, formylation, gamma-carboxylation, glycosylation, GPI anchor formation, hydroxylation, iodination, methylation, myristoylation, oxidation, pegylation, proteolytic processing, phosphorylation, prenylation, racemization, selenoylation, sulfation, transfer-RNA mediated addition of amino acids to proteins such as arginylation, and ubiquitination. (See, for instance, PROTEINS - STRUCTURE AND MOLECULAR PROPERTIES, 2nd Freeman and Company, New York (1993); Ed., T. E. Creighton, W. H. POSTTRANSLATIONAL COVALENT MODIFICATION OF PROTEINS, B. Johnson, Ed., Academic Press, New York, pgs. 1-12 (1983); Seifter et al., Meth. Enzymol. 182:626-646 (1990); Rattan et al., Ann. N.Y. Acad. Sci. 663:48-62 (1992)).

"SEQ ID NO:X" refers to a polynucleotide sequence described, for example, in Tables 1A, 1B or 2, while "SEQ ID NO:Y" refers to a polypeptide sequence described in column 11 of Table 1A and or column 6 of Table 1B. SEQ ID NO:X is identified by an integer specified in column 4 of Table 1B. The polypeptide sequence SEQ ID NO:Y is a translated open reading frame (ORF) encoded by polynucleotide SEQ ID NO:X. "Clone ID NO:Z" refers to a cDNA clone described in column 2 of Table 1A and/or 1B.

"A polypeptide having functional activity" refers to a polypeptide capable of displaying one or more known functional activities associated with a full-length (complete) protein. Such functional activities include, but are not limited to, biological activity, antigenicity [ability to bind (or compete with a polypeptide for binding) to an anti-polypeptide antibody], immunogenicity (ability to generate antibody which binds to a specific polypeptide of the invention), ability to form multimers with polypeptides of the invention, and ability to bind to a receptor or ligand for a polypeptide.

[79] The polypeptides of the invention can be assayed for functional activity (e.g. biological activity) using or routinely modifying assays known in the art, as well as assays described herein. Specifically, one of skill in the art may routinely assay secreted polypeptides (including fragments and variants) of the invention for activity using assays as described in the examples section below.

[80] "A polypeptide having biological activity" refers to a polypeptide exhibiting activity similar to, but not necessarily identical to, an activity of a polypeptide of the present invention, including mature forms, as measured in a particular biological assay, with or without dose dependency. In the case where dose dependency does exist, it need not be identical to that of the polypeptide, but rather substantially similar to the dose-dependence in a given activity as compared to the polypeptide of the present invention (i.e., the candidate polypeptide will exhibit greater activity or not more than about 25-fold less and, preferably, not more than about tenfold less activity, and most preferably, not more than about three-fold less activity relative to the polypeptide of the present invention).

## TABLE 1A

	Last	AA	of	ORF	141		9		18			78		25		41		139		30		20	
	First	AA of	Secreted	Portion	<i>L</i>				14			27		10		24		29		21		12	
Tact	AA	Jo	Sig	Pep	9				13			26		6		23		28		20		11	
Hiret I set	AA	of	Sig	Pep	1		1		1			1		-				1				1	
<b>▼</b>	SEQ		NO:	Y	515		516		517			518		519		520		521		522		523	
5° NT	First	AA of	Signal	Pep	83		1448		263			135		250		217		250		347		238	
	5' NT	Jo	Start	Codon			1448					135				217		250		347			
4, NL 3, NL	of	Clone Clone	Seq.		909		2377		1493			1300		888		3239		<i>L</i> 99		2318		330	
S, NT	of	Clone	Seq.		44		1275		1			1		1		1		1		1		1	
		Total	N	Seq.	509		2610		1493			1300		888		3239		<i>L</i> 99		2318		330	
ΤN	SEQ	А	S. S.	X	11		12		13			14		15		16		11		18		19	
				Vector	Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR			Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		pSport1		pSport1		pBluescript	
	ATCC	Deposit	No:Z	and Date	203917	04/08/99	203959	04/26/99	PTA-	793	09/27/99	203917	04/08/99	203979	04/29/99	203979	04/29/99	203917	04/08/99	203917	04/08/99	203979	04/29/99
			cDNA	Clone ID	H6BSF56	•	H6EDM64		H6EEC72			HACAB68		HACBJ56		HACBS22		HADDE71		HADDJ13		HADMB15	
			Gene	No.	_		2		3			4		5		9		7		8		6	

		Last	AA	Jo	ORF	21	17	55		11	54	01	13	38	94	359	38
		First	AA of	Secreted	Portion (	20	17	30			27			16	21	20	33
	Last	AA		Sig		19	16	29	1		56			15	70	19	32
	First Last	AA	of	Sig		1	-	-	1		-	-	-	-	_	-	1
	AA	SEQ	А	NO:	Y	524	525	526	1	527	528	529	530	531	532	533	534
5, NT	of	First	AA of	Signal	Pep	171	238	146	1	515	241	006	196	192	12	605	284
		S' NT	Jo	Start	Codon	171	238	146		515	241	006	196		12	605	284
	3' NT	of	Clone Clone	Seq.		743	1284	2890		785	874	2440	1346	1237	2345	2536	2182
	5' NT 3' NT	of		Seq.		-		100	,	-	-	843	-	-	Н	-	1
			Total	N	Seq.	743	1284	5684	1	785	874	2440	1346	1237	2345	2536	2182
	N	SEQ	О	ÿ N	Х	20	21	22	3	23	24	25	26	27	28	29	30
					Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR		Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0	pCMVSport
		ATCC	Deposit	No:Z	and Date	203917 04/08/99	203917 04/08/99	203917	2000110	203917 04/08/99	203979 04/29/99	203917 04/08/99	203979 04/29/99	203917 04/08/99	203959 04/26/99	PTA- 181 06/07/99	203917
				cDNA	Clone ID	HAGBQ12	HAGDW20	HAGEG10		нАСЕQ79	HAGFS57	HAGHIN57	HAHEA15	HAJAA47	HAJAY92	HAJBV67	HAJCH70
				Gene	No.	10	11	12	,	13	14	15	16	17	18	19	20

_								_																		
		Last	AA		ORF	1167		23		3		89		42		37		17		42		61		124		
		First	AA of		Portion	23		16				21		17		24				20		19		28		
	First Last	AA		Sig		22		15				20		16		23				19		18		27		
	_	AA		Sig	Pep	-		-		1				-		-		1		-		-		-		
	AA	SEQ	Ω	NO:	Y	535		536		537		538		539		540		541		542		543		544		
5' NT	Jo	First	AA of	Signal	Pep	8		250		262		18		268		296		271		93		521		17		
		5' NT	Jo	Start	Codon			250				18		268		296		271		93		521		17		
	3' NT	Jo	Clone	Seq.	1	4802		602		596		1380		903		1809		934		850		1596		720		
	5' NT 3' NT	of	Clone Clone	Seq.		7		1		1				1		95		1		1		293		1		
			Total	ZN	Seq.	5143		209		965		1380		903		1809		934		850		1713		720		
	N	SEQ	А	NO:	×	31		32		33		34		35		36		37		38		39	·	40		
					Vector	pSport1		Uni-ZAP XR		pSport1	•	pSport1														
		ATCC	Deposit	No:Z	and Date	203979	04/29/99	203917	04/08/99	203917	04/08/99	203917	04/08/99	203917	04/08/99	203917	04/08/99	203917	04/08/99	203917	04/08/99	203917	04/08/99	PTA-	794	09/27/99
				cDNA	Clone ID	HAOAG15		HAQAI92		HAQCE11		HATBI94		HATCB45		HATCD80		HATCI03		HATEH20		HBAGD86		HBCJL35		
				Gene	No.	21		22		23		24		25		56		27		28		29		30		

		ast	AA	of	ORF	124		T		[8	67	1	~		92			28		203		136		76		42	
		First L	AA of	_	Portion C	28		$\dagger$		t	07				6			17		25		16		25		17	
	Last	AA	ot		Pep	27		,	19	٤	61				∞			16		24		15		24		16	
	First	AA	of	Sig	Pep			]	-	,	<b>→</b>		_		-			1		-		_		1		1	
	AA	SEQ	А	NO:	Y	1015		1	545	,	546		547		548			549		550		551		552		553	
5' NT	Jo	First SEQ	AA of	Signal	Pep	1033			351	į	6/1		1016		550			110		28		1877		1036		586	
		5' NT	Jo	Start	Codon	1033			351	į	671									28		1877		9801			
		Jo	Clone	Seq.		1747			<b>687</b>	100,	1007		1829		802			069		1647		2392		1545		619	
	5' NT 3' NT	Jo	Clone Clone	Seq.		1027			<b>—</b>	+	320		764					1		_		1612		808		1	
	•		Total	Z	Seq.	2878			687	100	1007	1	1856		805			069		1647		2392		1782		619	
	Z	SEQ	П	NO:	X	511			41	Ţ	42		43		44			45		46		47		48		49	
					Vector	pSport1			pSport1		pSport1		Uni-ZAP XR		Uni-ZAP XR			Uni-ZAP XR									
		ATCC	Deposit	No:Z	and Date	PTA-	794	09/2/199	203917	77/00/10	203917	04/00/99	203917	04/08/99	PTA-	793	09/27/99	203917	04/08/99	203959	04/26/99	203917	04/08/99	203917	04/08/99	203917	04/08/99
				cDNA	$\overline{}$	HBCJL35			HBDAB91		HBDAB91		HBGBC29		HBGNC72			HBHAA05		HBHAA81		HBIAA59		HBIAC29		HBICW51	
				Gene	No.	30			31		32		33		34			35		36		37		38		39	

									5' NT					
	•			N		5' NT 3' NT	3, NT		Jo	AA	First	Last		
		ATCC		SEQ		Jo		5' NT	First	SEQ	AA	AA	First	Last
		Deposit		Ω	Total	Clone Clone	Clone	of	AA of	А		of	AA of	AA
Gene	cDNA	No:Z		NO:	N	Seq.	Seq.	Start	Signal	NO:			Secreted	of
No.	Clone ID	and Date	Vector	X	Seq.			Codon	Pep	Y	Pep	Pep	Portion	ORF
40	HBJAB02	203917	Uni-ZAP XR	20	1693		1665	84	84	554		27	78	34
		04/08/99												
41	HBJAC65	203917	Uni-ZAP XR	51	1685	-	892	137	137	555		13	4	23
		04/08/99												
42	HBJBM12	203917	Uni-ZAP XR	52	1135	-	1135	47	47	256				31
		04/08/99												
43	HBJCR46	203917	Uni-ZAP XR	53	3208	2270	3202	589	589	557	-		2	733
		04/08/99									.			
44	HBJDS79	203917	Uni-ZAP XR	54	2325	968	2325	1032	1032	558		37	38	107
		04/08/99												
45	HBJDW56	203917	Uni-ZAP XR	55	637		637		121	559				∞
		04/08/99												
46	HBJEL16	203979	Uni-ZAP XR	99	750		750	115	115	260	_	24	25	36
		04/59/99			•									
47	HBJFK45	203917	Uni-ZAP XR	57	543		543		430	561				∞
		04/08/69												
48	HBJIG20	PTA-	Uni-ZAP XR	28	637		637		321	562	-	16	17	77
		181												
		66/L0/90												
49	HBJKD16	203979	Uni-ZAP XR	59	1629	1	1629	78	78	563	_	18	19	31
		04/29/99												
20	HBMBM96	203917	pBluescript	09	1076	-	1076		170	564	_			4
		04/08/99												

								5, NT					
			NT		5' NT 3' NT	3, NT		Jo	AA	First Last	Last		
	ATCC		SEQ		Jo		5' NT	First SEQ	SEQ	AA	AA	First	Last
	Deposit		A	Total	Total Clone Clone	Clone	of	AA of	А	of	of	AA of	AA
cDNA	No:Z		SO.	N	Seq.	Seq.	Start	Signal NO:		Sig	Sig	Secreted	Jo
$\overline{}$	and Date	Vector	X	Seq.			Codon	Pep	Υ	Pep	Pep	Portion	ORF
HBMBX01	203917	pBluescript	19	1652	179	1458	363	363	595	_	18	19	78
	04/08/99								Ì				
HBMTM11	203917	Uni-ZAP XR	62	1639		1639	125	125	999	-	19	70	31
	04/08/99												
HBMTX26	203917	Uni-ZAP XR	63	1308	-	1308	107	107	267	-	46	47	68
	04/08/99												
HBMTY48	203917	Uni-ZAP XR	49	1891		1891	099	099	268	_	36	37	46
	04/08/66												
HBMUH74	PTA-	Uni-ZAP XR	9	726	_	726	344	344	569	<u></u>	13	14	78
	181												
	66/L0/90												
HBMWE61	203917	Uni-ZAP XR	99	1118	-	1118	238	238	570	-			6
HBNAX40	203917	Uni-ZAP XR	<i>L</i> 9	2793	2455	2793	2497	2497	571	1	18	19	49
	04/08/99												
HBNBJ76	203917	Uni-ZAP XR	89	1974	1469	1974		1603	572		53	30	89
	04/08/99												
HBQAB79	203917	Lambda ZAP	69	1331	1	1331	190	190	573				=
	04/08/99	П											
HBQAC57	203917	Lambda ZAP	70	2111		2111	146	146	574				53
	04/08/99	П											

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		Last	AA	of	ORF	48			16		-		35		9		254		23		32		32		55		55	
		First	AA of		Portion	28							18		13		18		16		14		6		27		16	
	Last	AA		Sig		27							17		12		17		15		13		∞		26		15	
	First	AA	of	Sig	Pep	_			-		1		1		1		-		-		1		-				-	
	AA	SEQ	А	ÖN:	≻	575			576		577		578		579		580		581		582		583		584		585	
5' NT	Jo	First	AA of	Signal NO:	Pep	447			119		1148		098		333		77		29		1588		995		177		131	
		5' NT	of	Start	Codon	447			119						333		LL								177		131	
	5' NT 3' NT	Jo	Clone Clone	Seq.		265		-	1010		1219		1392		813		1896		1276		1806		1732		1419		1052	
	5' NT	Jo	Clone	Seq.		129			41		1		628		1		1		1		1347		282		1		1	
			Total	N	Seq.	592			1010		1219		1392		813		9681		1276		2081		1732		1419		1052	
	K	SEQ	А	NO:	X	71			72		73		74		75		9/		77		78		79		80		81	
					Vector	Uni-ZAP XR			ZAP Express		ZAP Express		Uni-ZAP XR															
		ATCC	Deposit	No:Z	and Date	PTA-	181	66/10/90	203917	04/08/99	203917	04/08/99	203917	04/08/99	203917	04/08/99	203917	04/08/99	203979	04/29/99	203917	04/08/66	203917	04/08/99	203917	04/08/99		04/08/99
				cDNA	Clone ID	HBSAK32			HBXCM66		HBXCX15		HCDCY76		HCDDL48		HCE1G78		HCE2H52		HCE3B04		HCESF78		HCEDR26		HCEEE79	
				Gene	No.	19			62		63		49		65		99		29		89		69		70		71	

		st		<u>.</u>	H	8	T	\ <u>\</u>		J.,	-	l. <sub>c</sub>	Т		_				<u> </u>		<u> </u>			L	
		Last	AA	l of	ORF	23	43	265		15		25	٢	<del>-</del>	147		8		14		31			47	
		First	AA of	Secreted	Portion	16	31	17				17			24		32				20			26	
	Last	AA	of	Sig	Pep	15	30	16				16			23		31				19			25	
	First	AA	of	Sig	Pep	1		1		1		-	-	-	1		I.		-		1			1	
	AA	SEQ	А	NO:	Y	989	587	588		589		590	501	J71	592		593		594		595			596	
5' NT	Jo	First	AA of	Signal NO:	Pep	111	209	215		237		101	201	- 10C	304		539		101		1145			31	
		5' NT	of	Start	Codon		209	215		237		101							101		1145			31	
	3' NT	of	Clone Clone	Seq.		992	1229	1781		1305		1434	725	CC /	1359		2190		746		1633			1796	
	5' NT 3' NT	of		Seq.		-		4		1		1	-	<b>⊣</b>	62		334		1		1031			9//	
			Total	Z	Seq.	992	1229	1811		1305		1434	010	017	1359		2253		746		1728			1796	
	NŢ	SEQ	А	SON.	X	82	83	84		85		98	7.8	ò	88		68		96		91			92	
					Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR		Uni-ZAP XR		pSport1	nCnort 1	noded	pSport1		Uni-ZAP XR		Lambda ZAP	П	pBluescript			Lambda ZAP	П
		ATCC	Deposit	No:Z	and Date	203917 04/08/99	203917 04/08/99	203917	04/08/99	203917	04/08/99	203917 04/08/99	203017	04/08/99	203979	04/29/99	203917	04/08/99	203917	04/08/99	PTA-	181	66/L0/90	203917	04/08/99
				cDNA	Clone ID	нсее025	HCEEU18	HCEFZ82		HCEGX05		HCFLN88	HCEI TOO		HCHAB84		HCMSX51		HCNC011		HCNSD29			нсовн72	
				Gene	No.	72	73	74		75		92	77	:	78		79		08		81			82	

									5' NT					
				N		5' NT 3' NT	3' NT		of	AA	First Last	Last		
		ATCC		SEQ		Jo	Jo	5' NT	First	SEQ	AA	AA	First	Last
		Deposit		О	Total	Clone Clone	Clone	Jo	AA of		Jo	of	AA of	AA
Gene		No:Z		: : :	Z	Seq.	Seq.	Start	Signal	NO:	Sig	Sig	Secreted	of
No.	Clone ID	and Date	Vector	X	Seq.			Codon	Pep	Y	Pep	Pep	Portion	ORF
83	9622ОЭН	203979	Lambda ZAP	93	3166	632	1455	782	782	265	1	70	21	45
		04/29/99	П											
84	Э5ГЭОЭН	203917	Lambda ZAP	94	1287	1	1287		728	869	П			
		04/08/99	П											
85	HCQCM24	203979	Lambda ZAP	95	1929	909	1929	815	815	599	1			38
		04/29/99	П											:
98	HCRAY10	203917	Uni-ZAP XR	96	881	1	788		141	009	1	36	37	145
		04/08/99												
87	HCRBF72	203917	Uni-ZAP XR	16	1264	101	1142	191	161	601	1	1	2	211
		04/08/99												
88	HCRNF78	203917	pSport1	86	892	П	892	363	363	602	-	22	23	46
		04/08/6												
68	HCUAF85	203917	ZAP Express	66	265	1	297	230	230	603	1	23	24	122
		04/08/66												
06	HCUCF89	203917	ZAP Express	100	530	1	530	189	189	604	1	18	19	59
		04/08/99												
91	HCUCK44	203957	ZAP Express	101	1143	578	1136	869	869	909	1	30	31	99
		04/26/99												
92	HCUDD64	203917	ZAP Express	102	405	150	389	256	256	909	1	35	36	49
		04/08/99												
93	HCWAE64	203917	ZAP Express	103	471	_	471		410	209	1			5
		04/08/99												:

AA First Last SEQ AA AA AA DO: Sig Sig Y Pep										S' NT					
ATCC   APC   Deposit   D					N		5° NT	3, NT		of	AA	First	Last		
CDNA         No.Z         No.Z <th< td=""><td></td><td></td><td>ATCC</td><td></td><td>SEQ</td><td></td><td>of</td><td></td><td>5' NT</td><td>First</td><td>SEQ</td><td>AA</td><td>AA</td><td>First</td><td>Last</td></th<>			ATCC		SEQ		of		5' NT	First	SEQ	AA	AA	First	Last
CDNA         No.Z         No.Z         NO. No.Z         NO. No.Z         NO. No.Z         Seq.         Start         Signal No. Signal         NO. Signal         No. Sig			Deposit		А	Total	Clone	Clone		AA of	А	of	Jo	AA of	AA
Clone ID         and Date         Vector         X         Seq.         Codon         Pep         Y         Pep	Gene	cDNA	No:Z		NO:	Ĭ	Seq.	Seq.	Start	Signal	NO:			Secreted	of
HCWFU39         203917 (A)08/99 (A)08/99         ZAP Express (A)08/99 (A)08/99         104 (A)08/99 (A)08/99         105 (A)08/99 (A)08/99         106 (A)08/99 (A)08/99         107 (A)08/99 (A)08/99         107 (A)08/99 (A)08/99         108 (A)08/99 (A)08/99 <t< td=""><td>No.</td><td>Clone ID</td><td>and Date</td><td>Vector</td><td>X</td><td>Seq.</td><td></td><td>_</td><td>Codon</td><td></td><td>Y</td><td></td><td>Pep</td><td>Portion</td><td>ORF</td></t<>	No.	Clone ID	and Date	Vector	X	Seq.		_	Codon		Y		Pep	Portion	ORF
HCWUL09         203917 (A)08/99         LAP Express         105         761         3         761         333         333         609         1           HDHAA42         203917 (A)08/99         CMVSport         106         943         1         943         48         48         610         1         25           HDHAA42         203917 (A)08/99         CMVSport         107         497         1         497         48         48         610         1         25           HDHEB76         203917 (A)08/99         CMVSport         107         497         1         497         416         611         1         11           HDPCW16         203960 (A)26/99         DCMVSport         109         1550         1         1550         23         23         613         1         17           HDPD158         203960 (A)26/99         DCMVSport         110         1997         1         1997         279         614         1         19           HDPF1043         203960 (A)26/99         DCMVSport         111         2582         3         2582         186         615         1         19           HDFF1043         203960 (A)26/99         DCMVSport         112<	94	HCWFU39	203917	ZAP Express		467	1	467	282	282	809	-	6	10	22
HCWUL09         203917         ZAP Express         105         761         3         761         333         333         609         1           HDHAA42         203917         PCMVSport         106         943         1         943         48         48         610         1         25           HDHEB76         203917         PCMVSport         107         497         1         497         416         611         1         11           HDPCW16         203960         PCMVSport         108         1536         1         1536         172         172         612         1         38           HDPCW16         203960         PCMVSport         109         1550         1         1550         23         23         613         1         17           HDPDJ38         203609         PCMVSport         110         1997         1         1997         279         279         614         1         19           HDPFU43         203960         PCMVSport         111         2582         3         2582         186         16         1         19           40426/99         3.0         CMVSport         111         2582         3			04/08/99												_
HDHAA42         203917         pCMVSport         106         943         1         943         48         48         610         1         25           HDHEB76         203917         pCMVSport         107         497         1         497         416         611         1         11           HDPCW16         203960         pCMVSport         108         1536         1         1536         172         172         612         1         38           HDPCW16         203960         pCMVSport         109         1550         1         1550         23         23         613         1         17           HDPDI72         PTA-         pCMVSport         110         1997         1         1997         279         614         1         1           HDPFI10         PTA-         pCMVSport         111         2582         3         2582         186         615         1         19           HDPFU43         203960         pCMVSport         112         1904         1         1889         220         220         616         1         28           HDPFU43         203960         pCMVSport         112         1904         1	95		203917			761	3	761	333	333	609	1			=
HDHEB76 203917 pCMVSport 107 497 1 497 416 611 1 11 11	96	HDHAA42	203917	pCMVSport	106	943	1	943	48	48	610	-	25	26	26
HDHEB76         203917         pCMVSport         107         497         1         497         416         611         1           04/08/99         2.0         1         1536         172         172         612         1         18           HDPCW16         203960         pCMVSport         108         1550         1         1550         23         23         613         1         17           HDPD172         PTA-         pCMVSport         109         1550         1         1550         23         23         613         1         17           HDPD178         203960         pCMVSport         110         1997         1         1997         279         279         614         1         19           HDPFU43         203960         pCMVSport         111         2582         3         2582         186         186         615         1         19           HDPFU43         203960         pCMVSport         112         1904         1         1889         220         220         616         1         28           HDPFV18         203918         pCMVSport         112         1904         1         2187         161			04/08/99	2.0											
HDPCW16         203960         pCMVSport         108         1536         1         1536         172         172         612         1         38           HDPDI72         PTA-         pCMVSport         109         1550         1         1550         23         23         613         1         17           HDPDI72         PTA-         pCMVSport         110         1997         1         1997         279         279         614         1         17           HDPFF10         PTA-         pCMVSport         111         2582         3         2582         186         615         1         19           HDPFU43         203960         pCMVSport         112         1904         1         1889         220         20         616         1         28           HDPFU43         203960         pCMVSport         112         1904         1         1889         220         220         616         1         28           HDPFV18         203918         pCMVSport         113         2187         1         161         161         1           44/26/99         3.0         1         2187         1         2187         1         1	16	HDHEB76	203917	pCMVSport	107	497	1	497		416	611		=	12	12
HDPCW16         203960         pCMVSport         108         1536         1         1536         172         172         612         1         38           HDPDI72         PTA- 794         PCMVSport         109         1550         1         1550         23         23         613         1         17           HDPDI58         203960         PCMVSport         110         1997         1         1997         279         614         1         19           HDPFF10         PTA- 181         PCMVSport         111         2582         3         2582         186         615         1         19           HDPFU43         203960         PCMVSport         111         2582         3         2582         186         615         1         19           HDPFU43         203960         PCMVSport         112         1904         1         1889         220         220         616         1         28           HDPFV18         203960         PCMVSport         112         1904         1         1889         220         220         616         1         28           HDPFV18         203918         PCMVSport         113         2187 <t< td=""><td></td><td></td><td>04/08/99</td><td>2.0</td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td></t<>			04/08/99	2.0											
HDPDI72       PTA- PTA- PCMVSport 109       1550       1       1550       23       23       613       1       17         794       3.0       109/27/99       3.0       109/27/99       109/27/99       109/27/99       109/27/99       109/27/99       110       1997       1       1997       279       279       614       1       17         HDPFF10       PTA- PCMVSport 111       2582       3       2582       186       186       615       1       19         HDPFU43       203960       PCMVSport 112       1904       1       1889       220       616       1       28         HDPFY18       203918       PCMVSport 113       2187       1       2187       161       161       1       1       1	86	HDPCW16	203960	pCMVSport	108	1536	-	1536	172	172	612	_	38	39	55
HDPDI72         PTA-PTA-PTA-PTA-PTA-PTA-PTA-PTA-PTA-PTA-			04/26/99	3.0											
HDPDJ58       203960 pCMVSport 110       1997       1 1997       279       279       614       1         HDPFF10       PTA- pCMVSport 111       2582       3       2582       186       186       615       1       19         HDPFU43       203960 pCMVSport 112       1904       1       1889       220       220       616       1       28         HDPFV18       203918 pCMVSport 113       2187       1       2187       161       161       161       1       1	66	HDPDI72	PTA-	pCMVSport	109	1550	_	1550	23	23	613		17	18	120
HDPDJ58         203960 pCMVSport 110         1997 1 1997         279         279 614         1           HDPFF10         PTA- pCMVSport 111         2582         3 2582 186         186 615         1 199           HDPFU43         203960 pCMVSport 112         1904 1 1889 220         220 616         1 28           HDPFV18         203918 pCMVSport 113         2187 1 2187 161         161 617 1			794	3.0											
HDPDJ58         203960         pCMVSport         110         1997         1         1997         279         279         614         1           HDPFF10         PTA-         pCMVSport         111         2582         3         2582         186         186         615         1         19           181         3.0         181         3.0         1         1889         220         220         616         1         28           HDPFU43         203960         pCMVSport         112         1904         1         1889         220         220         616         1         28           HDPFY18         203918         pCMVSport         113         2187         1         2187         161         161         1         1			09/27/99												
HDPFF10       PTA-       pCMVSport       111       2582       3       2582       186       186       615       1       19         181       3.0       06/07/99       203960       pCMVSport       112       1904       1       1889       220       220       616       1       28         HDPFU43       203918       pCMVSport       113       2187       1       2187       161       161       161       1       1	100	HDPDJ58	203960	pCMVSport	110	1997	-	1997	279	279	614	_			70
HDPFF10 PTA- pCMVSport 111 2582 3 2582 186 186 615 1 19 19			04/26/99	3.0											
HDPFU43 203960 pCMVSport 112 1904 1 1889 220 220 616 1 28	101	HDPFF10	PTA-	pCMVSport	-	2582	$\mathcal{C}$	2582	186	186	615	_	19	70	425
HDPFU43 203960 pCMVSport 112 1904 1 1889 220 220 616 1 28 64/26/99 3.0 HDPFY18 203918 pCMVSport 113 2187 1 2187 161 161 617 1			181	3.0											
HDPFU43         203960         pCMVSport         112         1904         1         1889         220         220         616         1         28           04/26/99         3.0         3.0         1         2187         1         2187         161         161         617         1           04/08/99         3.0         3.0         3.0         1         2187         161         161         617         1			66/L0/90												
HDPFY18 203918 pCMVSport 113 2187 1 2187 161 161 04/08/99 3.0	102	HDPFU43	203960	pCMVSport	112	1904	1	1889	220	220	616	1	28	59	52
HDPFY18 203918 pCMVSport 113 2187 1 2187 161 161 161 04/08/99 3.0			04/26/99	3.0											
	103	HDPFY18	203918	pCMVSport	113	2187	П	2187	161	191	617	_			7
			04/08/66	3.0											

٠								5' NT					
			N		5' NT 3' NT	3' NT		Jo	AA	AA First Last	Last		
	ATCC		SEQ		Jo	of	5' NT	First	SEQ	AA	AA	First	Last
	Deposit		П	Total	Clone Clone	Clone	Jo	AA of			of	AA of	AA
cDNA	No:Z		NO:	N	Seq.	Seq.	Start	Signal NO:		Sig	Sig		of
Clone ID	and Date	Vector	X	Seq.			Codon	Pep	Y		Pep	E	ORF
HDPGE24	203960	pCMVSport	114	2625	1	2625	173	173	618	_	11	12	73
	04/26/99	3.0											
HDPIU94	203960	pCMVSport	115	2196	21	2196	208	208	619	_	21	22	23
	04/26/99	3.0											
HDPOC24	203960	pCMVSport	116	1777	302	1725	418	418	620	_	23	24	133
	04/26/99	3.0										1	
HDPOL37	203960	pCMVSport	117	1489	1	1489	189	189	621	-	32	33	62
	04/56/99	3.0											
HDP0076	203960	pCMVSport	118	645	1	645		109	622	-	15	16	16
	04/26/99	3.0											
<b>Н</b> DРРD93	203960	pCMVSport	119	701	1	701	28	78	623	_			12
	04/26/99	3.0											
HDPPQ30	203960	pCMVSport	120	1063	_	1063	220	220	624	-	22	23	38
	04/26/99	3.0											1
HDPPW82	203959	pCMVSport	121	552	_	552	395	395	625				29
	04/26/99	3.0											
HDPXN20	203960	pCMVSport	122	1756	-	1756	61	61	979		20	21	41
	04/26/99	3.0											
HDQHIM36	PTA-	pCMVSport	123	1547	_	1547	129	129	627		18	19	48
	181	3.0											
	66/L0/90												
HDTAU35	203960	pCMVSport	124	377	-	377	260	260	628	-	12	13	17
	04/26/99	2.0											

		Last	AA	of	ORF	33		2		53		9		7		7		52		13		36		42		41	
		First	AA of	Secreted	Portion	23		17		18		19						6				17		78		19	
	Last	AA	of	Sig	Pep	22		91		17		18						∞				16		27		18	
	First Last	AA	of	Sig	Pep			_		_		-		1		1		-		_		-		_		-	
	AA	SEQ	Д		Y	629		630		631		632		633		634		635		989		637		638		639	
S' NT	Jo	First	AA of	Signal NO:	Pep	191		164		375		345		360		1731		321		10		273		170		41	
		5' NT	Jo	Start	Codon	191		164				345		360		1731		321				273		770		41	
	3, NT		Clone	Seq.		099		829		2261		525		1663		3034		809		999		1569		1323		845	
	5' NT 3' NT	Jo	Clone Clone	Seq.		1		_		_		1		308		1679						236		638		1	
			Total	Ĭ	Seq.	099		8/9		2261		525		1663	-	3034		608		995		1569		1323		845	
	L	SEO	А	NO:	X	125		126		127		128		129		130		131		132		133		134		135	
					Vector	pCMVSport	2.0	pCMVSport	2.0	pCMVSport	2.0	pCMVSport	2.0	Uni-ZAP XR													
		ATCC	Deposit	No:Z	and Date	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	00/90/10
				cDNA	$\overline{}$	HDTAV54		HDTFX18		HDTGW48		HDTLM18		HE2CA60		HE2CA60		HE2CH58		HE2CM39		HE2HC60		HE2P093		HE6AU52	
				Gene	Š.	115		116		117		118		119		120		121		122		123		124		125	

		Last	AA	of	ORF	62		25		46		576		33		47		47		43		43		471		43	
		First I	AA of		Portion	11		21		15	$\neg$	27		19		34		34		76		76		7		56	
	Last	AA	of		Pep	10		20		14		56		18		33		33		25		25		П		25	
	First Last	AA	of	Sig	Pep	_		-		-		-				-		_		_		_		_		_	
	AA	SEQ	А	NO	Y	640		641		642		643		644		645		949		647		648		649		920	
5' NT	Jo	First	AA of	1	Pep	295		38		171		145		210		155		155		157		1074		2		2268	
		5' NT	oę	Start	Codon			38		171		145		210		155		155		157		1074				2268	
	3, NT		Clone	Seq.		1526		941		867		1994		1526		1887		1887		1978		2891		4890		4085	
	5' NT 3' NT	Jo	Clone Clone	Seq.		1						1				_		1		_		918		2918		2114	
		<u>=</u>	Total	Z	Seq.	1526		941		867		2000		1526		1887		1887		1995		2908		4907		4102	
	IN	SEQ	Έ	NO:	X	136		137		138		139		140		141		142		143		144		145		146	
					Vector	Uni-ZAP XR																					
		ATCC	Deposit	No:Z	and Date	203960	04/26/99	203960	04/26/99	203979	04/29/99	203979	04/29/99	203960	04/26/99	203979	04/29/99	203979	04/29/99	203979	04/29/99	203979	04/29/99	203979	04/29/99	203979	04/29/99
				cDNA	$\overline{}$	HE6CS65		HE6D092		HE6EY13		HE6FU11		HE6FV29		HE8FC45		HE8FC45		HE8FD92		HE8FD92		HE8FD92		HE8FD92	
				Gene	No.	126		127		128		129		130		131		132		133		134		135		136	

ATCC	_									5' NT					
ATCC					N		5' NT	3, NT		of		First	Last		
CDNA         No:Z         NO: NT         Seq. Seq. Seq. Seq. Start         Signal NO: Seq. Seq. Seq. Seq. Seq. Seq. Seq. Seq.			ATCC		SEQ		of	of	5' NT		SEQ		AA	First	Last
CDNA         No:Z         NO:Z         NO:Z         NO:Z         NO:Z         NO:Z         Seq.         NO:Z         NO:Z         Seq.         Seq.         Seq.         Seq.         Seq.         NO:Z         NO:Z <th< td=""><td></td><td></td><td>Deposit</td><td></td><td>О</td><td>Total</td><td>Clone</td><td>Clone</td><td>Jo</td><td>AA of</td><td></td><td></td><td>of</td><td>AA of</td><td>AA</td></th<>			Deposit		О	Total	Clone	Clone	Jo	AA of			of	AA of	AA
Clone ID         and Date         Vector         X         Seq.         Codon         Pep         Y         Pep           HE8FD92         203979         Uni-ZAP XR         147         3977         1986         3960         2141         2141         651         1           HE8C996         PTA-         Uni-ZAP XR         148         2036         1         2036         118         118         652         1           66/07/99         PTA-         Uni-ZAP XR         149         2204         1400         2204         1413         1413         653         1           HE8TY46         PTA-         Uni-ZAP XR         150         1047         47         1047         55         55         654         1           HE9CY05         203960         Uni-ZAP XR         151         2114         1         211         212         655         1           HEBCI18         203960         Uni-ZAP XR         153         1121         713         1050         855         855         657         1           HEBCI18         203960         Uni-ZAP XR         153         1121         713         1050         855         855         657         1	Gene	cDNA	No:Z		NO:			Seq.	Start	Signal	NO:		Sig	Secreted	of
HESFD92         203979 (April 2AP XR)         148 (April 2AP)         149 (April 2AP)         141 (April 2	No.	Clone ID	and Date	Vector	X	Seq.			Codon	Pep	Y		Pep	Portion	ORF
HE8SG96       PTA-       Uni-ZAP XR       148       2036       1       2036       118       118         06/07/99       181       PTA-       Uni-ZAP XR       149       2204       1400       2204       1413       118         HE8TY46       PTA-       Uni-ZAP XR       150       1047       47       1047       55       55         HE9CY05       203960       Uni-ZAP XR       151       2114       1       2111       212         HE9CG20       203960       Uni-ZAP XR       152       676       1       676       319       319         HEBCT18       203960       Uni-ZAP XR       153       1121       713       1050       855       855         HEBCY54       203960       Uni-ZAP XR       154       1189       1       1189       172       172         HEBDQ91       203960       Uni-ZAP XR       155       1820       1       1820       681       681         HEBDQ91       203960       Uni-ZAP XR       155       1820       1       1820       681       681         HEBDQ91       203960       Uni-ZAP XR       156       1573       1007       1573       1211	137	HE8FD92	203979	Uni-ZAP XR	147	268	1986	3960	2141	2141	651	1	25	79	43
HE8SG96         PTA- 181         Uni-ZAP XR         148         2036         1         2036         118         118           06/07/99 1838         181 1838         Ani-ZAP XR         149         2204         1400         2204         1413         1413           HE9CY05         203960 04/26/99         Uni-ZAP XR         150         1047         47         1047         55         55           HE9CG20         203960 04/26/99         Uni-ZAP XR         152         676         1         676         319         319           HEBCI18         203960 04/26/99         Uni-ZAP XR         153         1121         713         1050         855         855           HEBDF77         203960 04/26/99         Uni-ZAP XR         154         1189         1         1189         172         172           HEBDQ91         203960 04/26/99         Uni-ZAP XR         155         1820         1         1820         681         681           HEBDQ91         203960 04/26/99         Uni-ZAP XR         155         1820         1         1820         681         681           HEBDQ91         203960 04/26/99         Uni-ZAP XR         156         1573         1007         1573         1211 </td <td></td> <td></td> <td>04/29/99</td> <td></td>			04/29/99												
HESTY46 PTA- Uni-ZAP XR 149 2204 1400 2204 1413 1413 1413 (55/09/00)  HE9CY05 203960 Uni-ZAP XR 151 2114 1 2111 212 212 (64/26/99)  HE9GG20 203960 Uni-ZAP XR 152 676 1 676 319 319 (64/26/99)  HEBCY54 203960 Uni-ZAP XR 153 1121 713 1050 855 855 (64/26/99)  HEBCY54 203960 Uni-ZAP XR 154 1189 1 1189 172 172 (681 681 681) (64/26/99)  HEBDQ91 203960 Uni-ZAP XR 155 1820 1 1820 681 681 (681 681) (64/26/99)	138	HE8SG96	PTA-	Uni-ZAP XR	148	2036	1	2036	118	118	652	1	17	18	24
HESTY46 PTA- Uni-ZAP XR 149 2204 1400 2204 1413 1413 1413 1413 1413 1413 1413 14			181												
HE8TY46         PTA- 1838         Uni-ZAP XR 149         149         2204 1400         1413 1413         1413			06/20/90												
HE9CY05       203960	139	HE8TY46	PTA-	Uni-ZAP XR	149	2204	1400	2204	1413	1413	653	1	18	19	187
HE9CY05       203960       Uni-ZAP XR       150       1047       47       1047       55       55         HE9EA10       203960       Uni-ZAP XR       151       2114       1       2111       212         HE9GG20       203960       Uni-ZAP XR       152       676       1       676       319       319         HEBCI18       203960       Uni-ZAP XR       153       1121       713       1050       855       855         HEBCY54       203960       Uni-ZAP XR       154       1189       1       1189       172       172         HEBDF77       203960       Uni-ZAP XR       155       1820       1       1820       681       681         HEBDQ91       203960       Uni-ZAP XR       156       1573       1007       1573       1211         HEBDQ91       203960       Uni-ZAP XR       156       1573       1007       1573       1211	-		1838												
HE9CY05         203960         Uni-ZAP XR         150         1047         47         1047         55         55           HE9EA10         203960         Uni-ZAP XR         151         2114         1         2111         212           HE9GG20         203960         Uni-ZAP XR         152         676         1         676         319         319           HEBCI18         203960         Uni-ZAP XR         153         1121         713         1050         855         855           HEBCY54         203960         Uni-ZAP XR         154         1189         1         1189         172         172           HEBDF77         203960         Uni-ZAP XR         155         1820         1         1820         681         681           HEBDQ91         203960         Uni-ZAP XR         155         1820         1         1820         681         681           HEBDQ91         204/26/99         104/26/99         1         1573         1007         1573         1211			02/06/00												
04/26/99       104/26/99       203960       Uni-ZAP XR       151       2114       1       2111       212         04/26/99       04/26/99       152       676       1       676       319       319         203960       Uni-ZAP XR       153       1121       713       1050       855       855         04/26/99       04/26/99       1       189       1       1189       172       172         04/26/99       04/26/99       1       185       1820       1       1820       681       681         04/26/99       203960       Uni-ZAP XR       155       1820       1       1820       681       681         04/26/99       203960       Uni-ZAP XR       156       1573       1007       1573       1211	140	HE9CY05	203960	Uni-ZAP XR	150	1047	47	1047	55	25	654	1	21	22	235
HE9EA10         203960         Uni-ZAP XR         151         2114         1         2111         212           04/26/99         04/26/99         1         676         1         676         319         319           HEBCI18         203960         Uni-ZAP XR         153         1121         713         1050         855         855           HEBCY54         203960         Uni-ZAP XR         154         1189         1         1189         172         172           HEBDF77         203960         Uni-ZAP XR         155         1820         1         1820         681         681           HEBDQ91         203960         Uni-ZAP XR         155         1820         1         1820         681         681           HEBDQ91         203960         Uni-ZAP XR         156         1573         1007         1573         1211	$\neg$		04/26/99												
HE9GG20       203960       Uni-ZAP XR       152       676       1       676       319       319         HEBCI18       203960       Uni-ZAP XR       153       1121       713       1050       855       855         HEBCY54       203960       Uni-ZAP XR       154       1189       1       1189       172       172         HEBDF77       203960       Uni-ZAP XR       155       1820       1       1820       681       681         HEBDQ91       203960       Uni-ZAP XR       156       1573       1007       1573       1211		HE9EA10	203960	Uni-ZAP XR	151	2114	_	2111		212	655		28	29	78
HE9GG20         203960         Uni-ZAP XR         152         676         1         676         319         319           HEBCI18         203960         Uni-ZAP XR         153         1121         713         1050         855         855           HEBCY54         203960         Uni-ZAP XR         154         1189         1         1189         172         172           HEBDF77         203960         Uni-ZAP XR         155         1820         1         1820         681         681           HEBDQ91         203960         Uni-ZAP XR         156         1573         1007         1573         1211	$\neg$		04/26/99												
HEBCI18       203960 Uni-ZAP XR 153 1121       713 1050 855 855         HEBCY54       203960 Uni-ZAP XR 154 1189 1       1189 172 172         HEBDF77       203960 Uni-ZAP XR 155 1820 1       1820 681 681         HEBDQ91       203960 Uni-ZAP XR 155 1820 1       1820 681 681         HEBDQ91       203960 Uni-ZAP XR 156 1573 1007 1573       1211		HE9GG20	203960	Uni-ZAP XR	152	9/9	-	9/9	319	319	959	_			6
HEBC118         203960         Uni-ZAP XR         153         1121         713         1050         855         855           HEBCY54         203960         Uni-ZAP XR         154         1189         1         1189         172         172           HEBDF77         203960         Uni-ZAP XR         155         1820         1         1820         681         681           HEBDQ91         203960         Uni-ZAP XR         156         1573         1007         1573         1211			04/26/99												
HEBCY54       203960       Uni-ZAP XR       154       1189       1       1189       172       172         HEBDF77       203960       Uni-ZAP XR       155       1820       1       1820       681       681         HEBDQ91       203960       Uni-ZAP XR       156       1573       1007       1573       1211		HEBCI18	203960	Uni-ZAP XR	153	1121	713	1050	855	855	<i>L</i> 29	1	43	44	69
HEBCY54       203960       Uni-ZAP XR       154       1189       1       1189       172       172         04/26/99       Uni-ZAP XR       155       1820       1       1820       681       681         HEBDQ91       203960       Uni-ZAP XR       156       1573       1007       1573       1211			04/26/99												
HEBDF77       203960 04/26/99       Uni-ZAP XR 155 1820 1       1820 681 681       681         HEBDQ91       203960 04/26/99 04/26/99       Uni-ZAP XR 156 1573 1007 1573 1211       1211		HEBCY54	203960	Uni-ZAP XR	154	6811	1	1189	172	172	859	_	24	25	118
HEBDF77       203960       Uni-ZAP XR       155       1820       1       1820       681       681         04/26/99       Uni-ZAP XR       156       1573       1007       1573       1211			04/26/99												
04/26/99       List       1573       1007       1573       1211         04/26/99       HEBDQ91       1573       1007       1573       1211		HEBDF77	703960	Uni-ZAP XR	155	1820	1	1820	681	681	629	1	29	30	36
HEBDQ91   203960   Uni-ZAP XR   156   1573   1007   1573   1211			04/26/99												
04/26/99		HEBDQ91	203960	Uni-ZAP XR	156	1573	1007	1573		1211	099	1	56	30	41
			04/26/99												

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		Last	AA	Jo	ORF	29		42		34		20		17		22		22		10		51			15		
		First	AA of	Secreted	Portion	27		26		31						17		17				41			14		
	Last	AA	of	Sig	Pep	26		25		30						16		16				40			13		
	First Last	AA	of	Sig	Pep	1	·	1		1		1		1		1		1		1		-					
	AA	SEQ		NO:	Y	661		662		663		664		999		999		299		899		699			070		
5' NT	of	First	AA of	Signal NO:	Pep	200		106		59		215		82		147	:	147		440		154			664		
		5' NT	of	Start	Codon	200		106		59		215		82		147		147		440							
	3' NT	of	Clone Clone	Seq.		1304		1867		1125		2168		1260		1109		1109		1614		939			746		
	5' NT 3' NT	of	Clone	Seq.		1		1		1		1		1		12		12	w	204		1			1		
			Total	N	Seq.	1304		1867		1125		2168		1260		1109		1109		1614		939			746		
	NT	SEQ	А	NO:	X	157		158		159		160		161		162		163	·	164		165		•	166		
					Vector	Uni-ZAP XR		pSport1			Uni-ZAP XR																
		ATCC	Deposit	No:Z	and Date	203979	04/29/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99		04/29/99	PTA-	181	66/L0/90	PTA-	181	06/1/0/90
				cDNA	Clone ID	HEBFR46		HEBGE07		HEGAU15		HELAT35		HELBU54		HELGG84		HELGG84		HEMEY47		HEOMC46			HEPBA14		
				Gene	No.	147		148		149		150		151		152		153		154		155			156		

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		Last	AA	_	ORF	32		20	,	9		22		99		7		45		15		18		40		31	
		First	AA of	Secreted	Portion	27		19	8	87		17		14				18						25		23	
	Last	AA	Jo	Sig	Pep	26		18	r c	7.7		16		13				17						24		22	
	First Last	AA	Jo	Sig	Pep	1		-	,	_		1		1		. —				-		П		1		1	
	AA	SEQ	А	NO:	Y	671		672		673		674		912		9/9		<i>LL</i> 9		8/9		629		089		681	
5' NT	of	First SEQ	AA of	Signal	Pep	120		306	100	237		541		292		151		136		170		216		154		40	
		5' NT	of		Codon	150		306	į	237		541								170		216		154		40	
	3, NT	of	Clone	Seq.		1647		829	100	2285		1533		1778		871		887		1437		1205		1153		998	
	5' NT 3' NT	of	Total Clone Clone	Seq.		1		<b>-</b>	[	73		328		Ţ		-		1		1		1		1		1	
				Į	Seq.	1647		829	100	2285		1533		1778		871		887		1437		1205		1153		998	
	N	SEQ	П	NO:	X	167		891	,	169		170		1/1		172		173		174		5/1		9/1		<i>LL</i> 1	
					Vector	pCMVSport	3.0	pCMVSport	0.0	Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR	
		ATCC	Deposit	No:Z	and Date	203960	04/26/99	203960	04/20/99	203979	04/29/99	203979	04/29/99	203960	04/26/99	203979	04/29/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	_	04/26/99
				cDNA	$\overline{}$	<b>НЕ</b> QАН80		неов 189	- 1	HETCI16		HETDW58		HETEY67		HFCDW95		HFCEI04		HFCFD04		HFCFE20		HFEAY59		HFGAJ16	
				Gene	No.	157		158		159		160		191		162		163		164		165		166		167	

_		Last	AA	of	ORF	51	$\exists$		]	43	7	42		25		<del></del>	7	<del>ر</del> د دو		 23	١	45		30 30	Ţ	 9		7
					$\neg$		+		+		+		+		$\dashv$		ľ		+		$\dagger$		$\dashv$		+			_
		First	AA of	Secreted	Portion	22		<b>5</b> 8		23	,	18		17	1	25	- 1	25		24		78		19				
	Last	AA	Jo	Sig	Pep	21		27		22	ļ	17	,	16		24	1	78		23		27		18				
	First Last	AA	of	Sig	Pep			<del></del>				<del> </del>		_				<del>-</del>		_		_		_				
	AA	SEQ	А	:ON	X	682		683		684		685		989		<b>687</b>		889		689		069		691		692		
5' NT	of	First	AA of	Signal	Pep	700		175		283		243		9		9/9		414		2546		203		577		<i>L</i> 9		
		5' NT	of	Start	Codon	700		175		283		243		9		9/9		414		2546		203		577		<i>L</i> 9		
	3, NT		Clone	Seq.		1165		1275		1157		1885		1031		2735		2644		3114		1419		1941		820		
	5' NT 3' NT	of	Clone Clone	Seq.		454		110		_						341		-		2302				322				
			Total	K	Seq.	1280		1275		1157		1885		1031		2735		2644		3115		1419		1941		820		
	L	SEQ	П	ON	X	178		179		180		181		182		183		184		185		186		187		188		
					Vector	pSport1		pSport1		pSport1		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		
		ATCC	Deposit	No:Z	and Date	203960	04/26/99	203960	04/26/99	203979	04/29/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	PTA-	181	66/L0/90
			,	cDNA	$\sim$	HFIHZ75		HFIJA29		HFIJA68		HFKES05		HFKEU12		HFPCZ55		HFPDR62		HFPDS07		HFRAB10		HFTBM38		HFTDH56		
				Gene	No.	168		169		170		171		172		173		174		175		176		177		178		

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		Last	AA	ot		<u>~</u>	- 18	39	- 1	30		G 		- 1	97		o 	;	6I —	-	39	;	48	_	52	- 1		_
		First	AA of	Secreted	Portion			31		14	Ì	56		,	16		_		17	,	l9		33		15			
	Last	AA	Jo	Sig	Pep			30	,	13		25		,	15				16		<u>8</u>		35		14			
	AA First Last	AA		Sig	Pep	-		-				-			-		_		_				_		-		-	
	AA	SEQ		NO:	>	693		694		695		969			697		869		669		700		701		702		703	
5' NT	Jo	First SEQ	AA of	Signal	Pep	14		92		163		149			172		258		43		233		280		20		317	
		5' NT	Jo	Start	Codon			92				149			172		258		43		233							
	3, NT		Clone	Seq.		1236		1233		1520		1379			1001		1378		1316		1738		528		1054		1475	
	5' NT 3' NT	of	Clone Clone	Seq.		1		_		40		_			-		_		1		_		_		1		23	
		<u> </u>	Total	N	Seq.	1236		1233		1520		1379			1001		1378		1316		1738		528		1054		5061	
	LZ	SEO	<sup>′</sup> А	:ÖN	×	189		190		191		192			193		194		195		196		161		198		199	
				·	Vector	pBluescript		pBluescript		Lambda ZAP	П	Lambda ZAP	п		Lambda ZAP	П	Lambda ZAP	П	Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		pSport1	
		ATCC	Deposit	No:Z	and Date	203960	04/26/99	203960	04/26/99	203960	04/26/99	PTA-	181	66/L0/90	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99
				cDNA	_	2		HFVHW43		HFXAV37		HFXBN86			HFXBT66		HFXFZ46		HGBER72		HGBEY14		HGBGN34		HGBHP91		HGCAC19	
				Gene	No.	179		180		181		182		-	183		184		185		186		187		188		189	

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		Last	AA	of	ORF	6		6	n		36			4			41		223		47		24		<u>~</u>	
		First	AA of	Secreted	Portion						17			19				:	22		19		18			
,	Last	AA	of	Sig	Pep						16			18					21		18		17			
j	First Last	AA	of	Sig	Pep	_		1			<del></del>			-			-		_		_		1		_	
	AA	SEQ	А	NO:	X	704		705	902		707			708			402		710		711		712		713	
5' NT	of	First SEQ	AA of	ر ح	Pep	317		315	813		159			183			229		42		37		290		181	
		5' NT	of	Start	Codon						159			183			229		42		37		290		181	
	3, NT	Jo	Total Clone Clone	Seq.		1534		1473	1935		594			1589			1547		2632		1816		575		1584	
	5' NT 3' NT	Jo	Clone	Seq.		23		21	<i>L</i> 8		2			1			_		_				46		1	
			Total	N	Seq.	1534		1771	2014		594			1589			1547		2632		1816		575		1584	
	N	SEQ	П	NO:	X	200		201	202		203			204			205		206		207		808		506	
					Vector	pSport1		pSport1	port	3.0	pCMVSport	3.0		pCMVSport	3.0		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR	·	Lambda ZAP	П	Lambda ZAP	$\Pi$
		ATCC	Deposit	No:Z	and Date	203960	04/26/99	203960 04/26/99	203960	04/26/99	PTA-	181	66/L0/90	PTA-	793	09/27/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99
				cDNA	$\overline{}$	HGCAC19		HGCAC19	HHEAK45		HHEGS55			HHEOW19			HHFFF87		HHFFL34		HHFFS40		HHGCS78		HHGDT26	
				Gene	No.	190		191	192	-	193			194			195		196		197		198		199	

						_															-						$\neg$
	Last	AA	of	ORF	27		38		2	,	55		24			32		9		327		327		99		2	
	First	AA of	Secreted	Portion	19		53				18		15			11				30		30		98			
_	Last AA	Jo	Sig	Pep	18		28				11		14			10				29		56		35			
Д; #6	Filst		Sig	Pep	1		_		_		1		1			1		I		1		1		1		1	
~	SEO	<u>A</u>	NO:	Y	714		715		716		717		718			719		720		721		722		723		724	
5' NT	o. First		Signal	Pep	156		157		069		62		221			380		453		189		161		1239		19	
	5' NT		Start	Codon			157		-		62		221				·	453		189		191		1239		61	
2, NT	of Jo	Clone Clone	Seq.		1838		1147		803		1431		1277			531		1093		1976		1982		2142		1009	
5' NIT 3' NIT	of I	Clone	Seq.		1		-		27		1		1			1		1		151		153		1001		1	
		Total	N	Seq.	1838		1147		1049		1444		1277			531		1093		1980		1982		2154		1009	
Ę	SEO	<u>A</u>	NO:	X	210		211		212		213		214			215		216		217		218		219	·	220	
		,		Vector	Uni-ZAP XR		pBluescript		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR			Uni-ZAP XR		Uni-ZAP XR		pBluescript	SK-	pBluescript	SK-	pSport1		pBluescript	SK-
i	ATCC	Deposit	No:Z	and Date	203960	04/26/99	203960	04/26/99	203959	04/26/99	203917	04/08/99	PTA-	181	66/L0/90	203960	04/26/99	203960	04/26/99	203960	04/26/99	203960	04/26/99	203959	04/26/99	203957	04/26/99
			cDNA	Clone ID	HHPFU28		HHPSA85		HHSBI06		HHSBI65		HHSDI53			HHSFC09		HHSGL28		HILCA24		HILCA24		HISAT67		HJBCU75	
	-		Gene	No.	200		201		202		203		204			205		206		207		208		500		210	

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		Last	AA	of	ORF	6		27			34		18		43		117		281		50			20		36	
		First	AA of		Portion						23				22		11		21		15			14		19	
	Last	AA	of	Sig	Pep						22				21		10		20		14			13		18	
	First Last	AA	Jo	Sig		1		1			1		-		-		1		1		-	•		-		-	
	AA	SEQ	П	NO:	Y	725		726			727		728		729		730		731		732			733		734	
5' NT	Jo	First SEQ	AA of		Pep	527		207			2492		170		256		374		257		207			343		261	
		5' NT	Jo	Start	Codon	•		207											755		207			343		261	
	5' NT 3' NT	Jo	Clone Clone	Seq.		599		1017			2886		1298		686		628		1919		1181			1801		2007	
	5' NT	of	Clone	Seq.		1		1			2233		69		1		_		581					1		1	
			Total	Z	Seq.	599		101			2886		1298		686		628		6161		1181			1801		2007	
	N	SEQ	А	NO:	×	221		222			223		224		225		226		227		228			229		230	
					Vector	pCMVSport	3.0	pCMVSport	3.0		pCMVSport	3.0	Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		pCMVSport	2.0	pCMVSport	2.0		pCMVSport	2.0	pSport1	
		ATCC	Deposit	No:Z	and Date	203957	04/26/99	PTA-	181	66/L0/90	203959	04/26/99	203957	04/26/99	203957	04/26/99	203959	04/26/99	203959	04/26/99	PTA-	181	06/10/90	203957	04/26/99	203957	04/26/99
				cDNA	Clone ID	HJMAA03		HJMAV41			HJMAY90		HJPBE39		HJPBK28		HJPCH08		HKABU43		HKACI79			HKAFF50		HKGBF25	
				Gene	No.	211		212			213		214		215		216		217		218			219		220	

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Last		ot ORF	36	11	4	10	41	46	348	348	99	14	34
First	AA of	Secreted Portion	27				24	27	24	30	21		25
Last AA		Sig Pep	26				23	26	23	29	20		24
First AA	of G	Sig Pep	-	-	1	1	<b>–</b>	_		1	-	-	1
AA SEO	í A	N N	735	736	737	738	739	740	741	1016	742	743	744
5' NT of First	AA of	Signal Pep	572	214	390	172	184	550	66	75	522	45	25
S' NT		Start Codon	572	214		172	184	550	66	75	522	45	25
3' NT of	Clone	Seq.	750	1049	1098	797	652	1815	1488	1474	999	1842	1427
5' NT 3' NT of	Clone Clone	Sed.	343	-	-	-	1	425	-	163	254	12	1
	Total	Sea.	788	1049	1098	797	652	1815	1488	3179	721	1842	1427
NT SEO	<b>Д</b> ;	<u>;;</u> ×	231	232	233	234	235	236	237	512	238	239	240
		Vector	pBluescript	pBluescript	pBluescript	Uni-ZAP XR	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0	pCMVSport 3.0	Uni-ZAP XR	Uni-ZAP XR
ATCC	Deposit	No:Z and Date	203957 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99	PTA- 181 06/07/99	203979 04/29/99	203959 04/26/99	203959 04/26/99	203957 04/26/99	203957 04/26/99	203957 04/26/99
	j	cDNA Clone ID	HKIXC44	HKMLK03	НКМГМ95	HKTAB41	HLDBG17	HLDCA54	нгроп79	нгрои79	HLDRT09	HLHAP05	HLHCS23
	1	Gene No.	221	222	223	224	225	226	227	227	228	229	230

							71.4		5' NT		į			
		ATCC		SEO		of of		5' NT	oi First	SEQ	AA	AA	First	Last
		Deposit		<sup>′</sup> 白	Total	Total Clone Clone	Clone	of	AA of	А		of	AA of	AA
Gene	cDNA	No:Z		NO:		Seq.	Seq.	Start	Signal	: : :		_	Secreted	of t
		and Date	Vector	×	Seq.			Codon	Pep	X	Pep	Pep	Portion	ORF.
	HLB072	PTA-	pCMVSport 1	241	1768	1	1768	167	167	745	-	46	47	127
		792												
		06/12/160												,
232	HLICE88		pCMVSport 1	242	840	401	824		708	746	-			7
		04/26/99												
233	HLICO10	203957	pCMVSport 1	243	903	1	903	441	441	747	<del>-</del>	23	24	72
		04/26/99								İ				
234	HLJBS28	203957	pCMVSport 1	244	926	-	926	359	359	748	_			17
-		04/26/99												
235 F	HLMBW89	203957	Lambda ZAP	245	625	1	622	47	47	749	_	19	20	21
		04/26/99	П											
236	HLMGP50	203957	Lambda ZAP	246	1063		1063	214	214	750				9
		04/26/99	П											
237	HLMJB64	203957	Lambda ZAP	247	804	-	804	12	12	751	-	29	30	49
		04/26/99	П										,	- (
238 I	HLMMX62	203957	Lambda ZAP	248	768	-	268	185	185	752	_	17	18	78
	7	04/26/99	П											!
239	HLQAS12	PTA-	Lambda ZAP	249	2450	-	2450	305	305	753	_	Ξ	12	12
		793	п											
		09/27/99												
240	HLQCL64	PTA-	Lambda ZAP	250	2385	1652	2385		<u>س</u>	754	1	<b>—</b>	7	182
	•	181	п											
		66/L0/90												

		[ast	AA	ot	ORF	52		<del></del>	1	57		57		32			20		43		35		23		43		4	
		First 1		Sig Secreted	E	17		19				18		30			33		41		78		12		25			
	Last	AA	of	Sig	Pep	16		18		17		17		53			32		40		27		1		24			
	First Last	AA	of	Sig	Pep	_						-		<b>-</b>			_				1		_				-	
	AA	SEQ	А	S S	X	755		756		757		758		759			760		761		762		763		764		765	
5' NT	Jo	First	AA of	Signal	Pep	68		192		1751		220		700			122		81		95		363		258		1863	
		5' NT	of	Start	Codon	68		192		1751		220		200			122		81		95				258		1863	
	3' NT		Clone	Seq.		1243		2564		2488		947		2062			1716		788		1611		979		1146		2966	
	5' NT 3' NT	Jo	Clone Clone	Seq.		1				1542				1	-		_		1								1527	
			Total	IN	Seq.	1243		2564		2495		947		2062			1716		788		1611		979		1146		2967	
Γ	K	SEQ	A	NO:	X	251		252		253		254		255			256		257		258		259		260		261	
				****	Vector	Lambda ZAP	П	pCMVSport	3.0	pCMVSport	3.0	pCMVSport	3.0	pCMVSport	3.0		pCMVSport	3.0	pCMVSport	3.0	pCMVSport	3.0	pSport1		pSport1		pSport1	
		ATCC	Deposit	No:Z	and Date	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	PTA-	795	09/27/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203959	04/26/99
				cDNA	$\overline{}$	9		HLWAF06		HLWAU42		HLWAU42		HLWAV47			HLWBB73		HLWCN37		HLWDB73		HLYDF73		HLYEU59		HLYGB19	
				Gene	No.	241		242		243		244		245			246		247		248		249		250		251	

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		Last	AA	_	ORF!	73		42	450		35		450	_	35	$\perp$	35				56	$\dashv$	<b>∞</b>		34	_
		First	AA of	• 1	Portion	18		21	26		21		56		21		21		18		18				70	
	Last	AA	of		Pep	17		70	25		20		25		20		20		17		17				19	
	First Last	AA	of	Sig	Pep	—		_	-	l	1				1		-				_		_			
	AA	SEQ	А		X	99/		792	768		692		770		771		772		773		774		775		9//	
5' NT	Jo	First	AA of	Signal NO:	Pep	406		211	106		497		106		498		97		211		62		135		30	
		5' NT	of	Start	Codon	406		211	106	)	497		106		498		26		211		26		135		30	
	3, NT	of	Clone	Sed.		752		640	1670	)	1670		1670		1671		1301		443		1190		1204		2641	
	5' NT 3' NT	Jo	Clone Clone	Seq.		1		-	405	2	405		406		407		1		П		-	,	_		1	
			Total	Z	Seq.	752		640	1733		1733		1733		1735		1301		443		1190		1204		2641	
	N	SEQ	, П	Ö	X	262		263	26.4	}	265		266		267		268		269		270		271		272	
					Vector	pSport1		pSport1	IIn: 7AD VD		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Lambda ZAP	П
	•	ATCC	Deposit	No:Z	and Date	203957	04/26/99	203957	202017	04/08/99	203917	04/08/99	203917	04/08/6	203917	04/08/99	203917	04/08/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99
				cDNA	$\overline{}$	HLYGE16		HLYGY91	TD/40 A 704		HMCAZ04		HMCAZ04		HMCAZ04		HMCAZ04		HMCFH60		HMDAB29		HMDAD44		HMEBB82	
				Gene	No.	252		253	130	t C7	255		256		257		258		259		260		261		262	

		Last	AA	of	SRF F	33		17		93		31		9		32		 90 90		10		27		35		381	
		First	AA of	=	티	17		13		19		73				=		17								2	
	Last	AA	of		Pep	16		12		18		22				01	Ì	16						25		-	
	First	AA	Jo		Pep	1		_						-		-		_		_		_		_		<b>-</b>	ļ
	AA	SEQ	А	NO:	×	111		778		779		780		781		782		783		784		785		98/		787	
5' NT	of	First	AA of	Signal	Pep	006		622		113		195		229		1149		249		273		295		120		107	-
		5' NT	Jo	Start	Codon	006				113		195		229				249		273		295		120		107	
	3, NT	Jo	Clone	Seq.		2806		2219		1607		1064		1738		1772		2048		799		1396		2945		1667	
	5' NT 3' NT	Jo	Clone Clone	Seq.		884		362		П		_		-		-				_				1		442	
			Total	Ž	Seq.	2836		2276		1607		1064		1738		1772		2048		799		1396		2945		1667	
	Z	SEO	<sup>′</sup> Д	NO:	X	273		274		275		276		277		278		279		280		281		282		283	
					Vector	Lambda ZAP	П	Lambda ZAP	П	Lambda ZAP	П	Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		pSport1		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR	
		ATCC	Deposit	No:Z	and Date	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203979	04/29/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203959	04/26/99
				cDNA	Clone ID	HMEDE24		HMEDI90		HMELM75		HMIAK10		HMIBF07		HMICI80		HMICP65		HMJAK70		HMSBE04		HMSCL38		HIMSCR69	
				Gene	No.	263		264		265		566		267		268		569		270		271		272		273	

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		Last	AA	Jo	ORF	93	113	35		,	4 8	30	3	227		9		25		397		38		23	
		First	AA of	Secreted	Portion	21	25	12		6	$\mathcal{C}_{2}$	21	17	26		33		22		2		31		22	
	Last	AA	of	Sig	Pep	20	24	11		3	77	20	3	25		32		21		-		30		21	
j	First	AA	of	Sig	Pep	1	1	-		-	_	-	4	-				1		-		-	-	-	
	AA	SEQ	А	NO:	Y	788	789	790		5	/91	707	1	793		794		795		961		797		862	
5' NT	oţ	First	AA of	Signal	Pep	37	50	959		200	60/	710	2	239		1437		274		137		1015		288	
		5' NT	of	Start	Codon	28	50			9,1	60/	710	21	239		1437		274		137		1015		288	
	3' NT	of	Clone	Seq.		1724	2249	2205		0000	3839	0000	2007	2709		2546		1351		2596		2101		1224	
	5, NT $ 3$ , NT	Jo	Clone Clone	Seq.		1	-	1		-	-	099		-		1327		1		08		927	-	1	
			Total	Ľ	Seq.	1724	2249	2205		0000	2839	2000	2	2709		2556		1351		2596		2288		1224	
	Z	SEQ	О	NO:	X	284	285	286		000	/ 97	288	2	289		290		291		292		293		294	
					Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR		1 - 0/1/10	pcivi v sport 3.0	2	3.0	pCMVSport	3.0	pSport1		pSport1		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR	
		ATCC	Deposit	No:Z	and Date	203957	203979	PTA-	793	020000	04/29/99	203057	04/26/99	203918	04/08/99	203957	04/26/99	203979	04/29/99	203957	04/26/99	203957	04/26/99	203957	04/26/99
				cDNA	Clone ID	HMSHC86	HMSHU20	HMSHY25		TTA (TT ) TT	HM1AB//	HMITAE26		HMUAN45		HMVBC31		HMVDU15		HIMWBL03	:	HMWJF53		HNEAK81	
				Gene	No.	274	275	276		777	117	278	2	279		280		281		282		283		284	

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	Last	AA	of	ORF	34		58		33		32		43		30		32		114		46		59		243		
	First	AA of	Secreted	Portion	24		21		20		23		18		21		24		28		18		23		31		
Last	AA	Jo	Sig	Pep	23		20		19		22		17		20		23		27		17		22		30	-	
First Last	AA	Jo	Sig	Pep	1		1		1		I		1		1		1		1		1		1		1		
AA	• 1		Ö.	Y	661		800		801		802		803		804		805		908		807		808		809		
5' NT of	First	AA of	Signal	Pep	472		316		02		9/9		314		178		248		89		47		205		237		
	5' NT	Jo	Start	Codon	472		316		20		9/9				178		248								237		
3' NT	of	Clone Clone	Seq.	i	2710		463		2073		1442		1436		728		915		1156		989		1045		1425		
5' NT 3' NT	of	Clone	Seq.		225		-		1		428		1		1		1				1		-		1		
		Total	K	Seq.	2710		489		2073		1442		1436		728		915		1156		989		1045		1425		
IN	SEQ	А	SO.	X	295		296		297		298	·	299		300		301		302		303		304		305		
				Vector	Uni-ZAP XR		Uni-ZAP XR																				
	ATCC	Deposit	No:Z	and Date	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	PTA-	181	06/1/0/90
			cDNA	Clone ID	HNECL22		HNECW49		HNEDH88		HNFAC50		HNFGR08		HNFHF34		HNGAK51		HNGAM58		HNGBH53		HNGDQ38		HNGDX18		
			Gene	No.	285		286		287		288		586		290		291		262		293		294		295		

	Last	AA	ot	ORF	132			17		4		77		32		89		6			∞	í	7.3	•		45
	First	AA of	Secreted	Portion	19					25		22				16						ţ	47			18
Last	AA			Pep	18					77		21				15						ì	46			17
AA First Last	AA	of	Sig	Pep	_					_		1		-		-						ŀ	_			-
AA	SEQ			Y	1017			810		811		812		813	·	814		815			816	1	817			818
5' NT of	First	AA of	Signal NO:	Pep	231			73		58		30		184		181		25			221		252			415
	5' NT	of	Start	Codon	231									184		181							252			415
3, NT		Clone	Seq.		1411			1002		1103		1029		585		541		1195			1047		1246			1048
5' NT 3' NT	of	Clone Clone	Seq.		1			$\vdash$		П								1			1		<b>—</b>			1
		Total	K	Seq.	1411			1002		1103		1029		585		541		1195			1047		1246			1048
Ę	SEQ	А	NO:	X	513			306		307		308		309		310		311			312		313			314
				Vector	Uni-ZAP XR			Uni-ZAP XR			Uni-ZAP XR		Uni-ZAP XR		:	Uni-ZAP XR										
	ATCC	Deposit	No:Z	and Date	PTA-	181	06//0/90	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	203957	04/26/99	PTA-	795	09/27/60	203957	04/26/99	PTA-	181	66/L0/90	203959 04/26/99
			cDNA	Clone ID	HNGDX18			HNGDY34		HNGEA34		HNGEQ75		HNGGA68		HNGGP65		HNGHZ69			HNGIV64		HNGJB41			HNGKT41
			Gene	No.	295			296		297		298		299		300		301			302		303			304

				ΤN		5. NT 3. NT	3, NT		5' NT of	AA	First	Last		
		ATCC		SEQ		Jo		5' NT		•	AA	AA	First	Last
		Deposit		А	Total	Clone Clone	Clone	Jo	AA of	А	Jo	Jo	AA of	AA
Gene	cDNA	No:Z		NO:	Z	Seq.	Seq.	Start	Signal		Sig	_	Secreted	ot
No.	)	and Date	Vector	X	Seq.			Codon	Pep	Y	Pep	Pep	Portion	ORF
305	HNGMW45	203959	Uni-ZAP XR	315	1530	1	1530	452	452	819		56	27	43
		04/26/99												
306	HINGNK44	203959	Uni-ZAP XR	316	1178	302	1178	611	611	820		18	19	74
		04/26/99												
307	HINGNO53	203959	Uni-ZAP XR	317	825	_	825	467	467	821	-	15	16	34
		04/26/99												
308	HNGPJ25	203959	Uni-ZAP XR	318	853	129	853	544	544	822		20	21	25
		04/26/99					-							
309	HNHEN82	203918	Uni-ZAP XR	319	573	-	573		78	823	-	13	41	17
		04/08/99												
310	HINHFE71	203959	Uni-ZAP XR	320	903	-	903	298	298	824	_			21
		04/26/99												
311	HNHGK22	203918	Uni-ZAP XR	321	606	<del>-</del>	606	239	239	825	_	76	27	4
		04/08/99												
312	HNHHB10	203959	Uni-ZAP XR	322	901	-	901	215	215	826	-	28	53	29
		04/26/99												
313	HNHKS19	203959	Uni-ZAP XR	323	790	-	790	192	192	827	_	26	27	41
		04/26/99												
314	HNTBT17	PTA-	pCMVSport	324	1959		1959	91	91	828	_			9
		181	3.0											
		06/0/90												
315	HNTMH79	203959	pSport1	325	922	-	922	48	<del></del>	829	-	35	36	38
		1770711												

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		Last	AA	of	ORF	123		124		123		43		27		33		10		43		71		39		51		
		First	AA of	Secreted	Portion	70		20		77		36		∞		18				40		56		25		53		
	Last	AA			Pep	19		19		21		35		7		17				39		25		24		78		
	First	AA	of	Sig	Pep	П		_		-				_		_		-						_				
	AA	SEQ		_	X	830		831		832		833		834		835		836		837		838		839		840		
5' NT	of	First	AA of	_	Pep	148		148		778		43		173		101		248		1714		48		138		230		
		5' NT	of	Start	Codon					778		43				101				1714				138		230		
	3, NT	of	Clone	Seq.		068		892		1249		900		604		1117		927		2216		1356		1036		1349		
	5' NT 3' NT	Jo	$\circ$	Seq.		1		T		772	·	-		1		_		1		1449		П		1		1		
			Total	ZZ	Seq.	927		676		1298		006		604		1119		927		2218		1356		1036		1365		
Γ	K	SEQ	白	NO:	X	326		327		328		329		330		331		332		333		334		335		336		
					Vector	Uni-ZAP XR		pSport1		pCMVSport	2.0	pCMVSport	2.0															
		ATCC	Deposit	No:Z	and Date	203959	04/26/99	203959	04/26/99	203959	04/26/99	203918	04/08/99	203918	04/08/99	203917	04/08/99	203959	04/26/99	203959	04/26/99	203959	04/26/99	203959	04/26/99	PTA-	795	09/27/99
				cDNA	$\overline{}$	HOABP31		HOABP31		HOACG07		HODAG07		HODBB70		HODBV05		HODCZ32		HOEBK60		HOFAA78		HOFNB74		HOFNU55		
				Gene	No.	316		317		318		319		320		321		322		323		324		325		326		

		st	- ✓	٠	出					∞	,	<u></u>									I				1,0	
		Last	AA	Jo 1	ORF	20		39		198		28			24		83		28		33		88		65	
		First	AA of	Secreted	Portion	11		20		26					19		23		22		28		7		28	
	Last	AA	of	Sig	Pep	10		19		25					18		22		21		27		9		27	
	First	AA	of	Sig	Pep	1		1		1		1			1		1		1		1		1		1	
	AA	SEQ	А	NO:	Y	841		842		843		844			845		846		847		848		849		850	
5' NT	Jo	First	AA of	Signal	Pep	309		21		183		88			17		203		848		200		70		170	
		5' NT	of	Start	Codon	309				183		88			17		203		848		200				170	
	3, NT	Jo	Clone	Seq.	ı	1478		1125		1157		902			1552		897		1750	-	1129		1282		1911	
	5' NT 3' NT	of	Clone Clone	Seq.	•	1		1		1		1					1		622		1		-		1	
			Total	N	Seq.	1478		1125		1157		905			1552		268		1767		1129		1284		1161	
	Z	SEQ		NO:	X	337		338		339		340			341		342		343		344		345		346	
					Vector	pCMVSport	2.0	Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR			Uni-ZAP XR											
		ATCC	Deposit	No:Z	and Date	816602	04/08/99	656607	04/26/99	503959	04/26/99	PTA-	181	66/L0/90	203959	04/26/99	203918	04/08/99	203979	04/29/99	203918	04/08/99	203918	04/08/99	203959	04/26/99
				cDNA	Clone ID	HOGBF01		HORBS82		HORBV76		5LOGSOH			HOSEC25		HOSEI81		HOSEJ94		HOUCA21		HOUDE92		HOUDR07	
				Gene	No.	327		328		329		330			331		332		333		334		335		336	

Last	of ORF	<u> </u>	72	20	26	25	30	93	45	19	17
						-	(6)	100	4		
First AA of	<u> </u>		43		70		24	41	40	18	
Last AA of	Sig Pep		42		19	:	23	13	39	17	
First AA of	• <u> </u>	1	1	1	-	_	1	1	1	1	1
AA SEQ ID	NO: Y	851	852	853	854	855	856	857	858	859	860
5' NT of First AA of	Signal Pep	144	520	188	252	184	1021	188	258	94	38
5' NT of	Start Codon		520	188	252	184	1021	188	258	94	38
3' NT of Clone	Seq.	799	2882	1102	1129	2587	3097	582	835	628	352
5' NT 3' NT of of Otatal Clone Clone	Seq.	9/	457	45	-	-	803	-	-	_	1
Total	NT Seq.	833	2927	1249	1129	2587	3097	582	835	628	352
NT SEQ ID	×	347	348	349	350	351	352	353	354	355	356
	Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	pSport1	Uni-ZAP XR	Uni-ZAP XR				
ATCC Deposit	No:Z and Date	PTA- 181 06/07/99	203959	PTA- 793 09/27/99	203918 04/08/99	203918 04/08/99	203917 04/08/99	203959 04/26/99	203959 04/26/99	PTA- 181 06/07/99	PTA- 181 06/07/99
	cDNA Clone ID	HOUED72	HOUFS04	ноингг	HOVBD85	HPCAB41	HPCAL26	HPEAD23	HPFBA54	HPFCI36	HPFDI37
	Gene No.	337	338	339	340	341	342	343	344	345	346

				Į		5' NT 3' NT	3, NT		5' NT	4 4	Firet I act	I act		
		ATCC		SEQ		of In		5' NT	ب	SEQ		AA	First	Last
		Deposit		А	Total	Clone Clone	Clone	o	AA of	А			AA of	AA
Gene	cDNA	No:Z		:ON	ž	Seq.	Seq.		Signal NO:	NO:			Secreted	of
No.	Clone ID	and Date	Vector	X	Seq.			Codon	Pep	Y	Pep	Pep	Portion	ORF
347	HPIAA80	203959	Uni-ZAP XR	357	616	312	919		314	861	_	13	14	37
		04/26/99												
348	HPJBJ51	203959	Uni-ZAP XR	358	2793	522	2421	715	715	862	_	14	15	69
		04/26/99												
349	HPJBJ51	203959	Uni-ZAP XR	329	2795	523	2422	716	716	863	-	14	15	69
		04/26/99												
350	HPJBU43	PTA-	Uni-ZAP XR	360	575		575		242	864				17
		181												
		06/01/90												
351	HPJCW58	203918	Uni-ZAP XR	361	1165	1	1165	177	177	865	_	19	70	28
		04/08/99												
352	HPMBX22	203959	Uni-ZAP XR	362	454	-	454		211	998	-			19
		04/26/99												
353	HPMCJ84	203918	Uni-ZAP XR	363	788	-	788	83	83	867	-	22	23	38
	i	04/08/99		`										
354	HPMCV30	203918	Uni-ZAP XR	364	806		806	52	52	898		27	78	47
		04/08/99												
355	HPMFH77	203918	Uni-ZAP XR	365	1891	_	1891		251	698	_	11	12	35
		04/08/99												
356	HPQAX38	203979	Lambda ZAP	366	1157	41	1157		295	870		10	11	16
		04/23/99	П		,									
357	HPQAX38	203979	Lamb	367	1158	41	1158		295	871	_	10	11	16
		04/29/99	11											

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		Last	AA		ORF	34		35	_	134		36	_	35			8 	$\perp$	13		26		25			4	4
		First	AA of		Portion	31		34		19		23		17			17						8			25	
	Last	AA	of		Pep	30		33		18		22		16			16						19			24	
	First Last	AA	of	Sig	Pep	_		-		-		_		_			_		-		_		_			-	
	AA	SEQ	О	NO:	×	872		873		874		875		876			877		878		879		880			881	
2, NL	Jo	First	AA of	Signal NO:	Pep	85		16		684		1810		265			885		194		405		220			122	
		5' NT	of	Start	Codon	85		16		684		1810					885		194		405						
	3, NT		Clone	Seq.		2267		434		1648		2757		689			1680		325		854		1496			1135	
	5' NT 3' NT	Jo	Clone Clone	Seq.		_		_		558		1701		1			859		_		240		1			П	
			Total	Ņ	Seq.	2267		434		1673		2805		60 <i>L</i>			1760		325		878		1496			1135	
	Z	SEQ	<u>A</u>	NO:	X	368		369		370		371		372			373		374		375		376			377	
					Vector	Lambda ZAP	П	Lambda ZAP	П	Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR			pBluescript		Uni-ZAP XR		Uni-ZAP XR		pCMVSport	3.0		pCMVSport	3.0
		ATCC	Deposit	No:Z	and Date	203918	04/08/99	203918	04/08/99	203959	04/26/99	203959	04/26/99	PTA-	181	66/10/90	203959	04/26/99	203918	04/08/66	203959	04/26/99	PTA-	181	66/L0/90	203959	04/26/99
				cDNA	$\overline{}$	HPQCB83		HPQCC53		HPRBH85		HPRCA64		HPRCD35			HPTRM02		HPWBA29		HPWDK06		HRAAD30			HRADA42	
				Gene	No.	358		359		360		361		362			363		364		365		366			367	

					IL.			$\neg$				1		$\neg$		Т		-		7				П			$\neg$
		Last	AA		ORF	253			65		63		31		46		6		72		10		10		33		
		First	AA of		Portion	40			18		53		28		18				21						56		
	Last	AA	of	Sig	Pep	39			17		28		27		17				20						25		
	First Last	AA	of	Sig	Pep				-		1		_		-		_				_		-		-		
	AA	SEQ	А	0 N	Y	882			883		884		885		988		887		888		688		068		891		
5' NT	Jo	First	AA of	Signal NO:	Pep	691			198		233		578		215		32		649		120		1958		20		
		5' NT	of	Start	Codon	169			198		233		578						649		120				20		
	3, NT	of	Clone	Seq.		2684			1206		1324		1500		773		543		1638		726		2271		281		
	5' NT 3' NT	of	Total Clone Clone	Seq.		1	•		17		1		547		1		1		711				1687		1	"	
				N	Seq.	2704			1225		1324		1500		9//		543		1681		728		2301		281		
	N	SEQ	Ω	SO.	X	378			379		380		381		382		383		384		385		386		288		
					Vector	pCMVSport	3.0		pCMVSport	3.0	pCMVSport.		Uni-ZAP XR	,	Uni-ZAP XR												
		ATCC	Deposit	No:Z	and Date	PTA-	181	66/L0/90	203959	04/26/99	203959	04/26/99	203918	04/08/99	203959	04/26/99	203959	04/26/99	203959	04/26/99	203959	04/26/99	203959	04/26/99	PTA-	181	66/120/90
				cDNA	$\overline{}$	HRADF49			HRADN25		HRADT25		HRDAI17		нкррозэ		HRDER22		HRDEX93		HRDFK37		HRGBD54		HROEA08		
				Gene	No.	368			369		370		371		372		373		374		375		376		377		

	Last	AA	of	ORF	56		22		63		37		98		36		10		13		16		12		0/	
	First	AA of	Secreted	Portion	81		<i>L</i> 1		30				<i>L</i> 1		91		8				20		13		17	
Last	AA	of	Sig	Pep	17		16		29			,	91		15		L				19		11		70	
First Last	AA	of	Sig	Pep	1		1		1		1		1		1		1		1		1		1		1	
AA	SEQ	П	NO:	Υ	892		893		894		895		968		268		868		668		006		901		905	
5° NT of	First	AA of	Signal	Pep	99		129		159		124		106		445		129		304		257		473		647	
	5, NT	Jo	Start	Codon			129				124		901		445				304		257				<i>L</i> 49	
3' NT	Jo	Clone Clone	Seq.		1001		262		349		6101		214		554		1213		882		1648	!	762		1474	
5' NT 3' NT	Jo	Clone	Seq.		1		1		1		1		1		1		1		I		1		1		452	
		Total	NT	Seq.	1061		595		349		1019		214		554		1273		882		1648		762		1474	
NT	SEQ	Ω	NO:	X	388		389		390		391		392		393		394		395		396		397		398	
				Vector	Uni-ZAP XR		pBluescript		Uni-ZAP XR																	
	ATCC	Deposit	No:Z	and Date	203918	04/08/99	203959	04/20/99	203959	04/26/99	503959	04/26/99	503959	04/26/99	503959	04/26/99	203959	04/26/99	656807	04/26/99	203918	04/08/99	203918	04/08/99	626807	04/29/99
			cDNA	Clone ID	HSAVA08		HSAVW42		HSAWN53		HSAWZ40		HSAYC41		HSDZM54		HSHBF76		HSIFG47		HSJBY32		HSKDR27		HSLHG78	
			Gene	No.	378		379		380		381		382		383		384		385		386		387		388	

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		Last	AA	Jo	ORF	41		4		14		62		42		99		Ξ			265		21			21	
ı		First	AA of			21						15		12		29					15						
	Last	AA	of	Sig	Pep	20						14		11		28					14						
	AA First Last	AA	of	Sig	Pep			1		-		1				1		-			-		-			1	
	AA	SEQ	白			903		904		905		906		706		806		606			910		911		-	912	
5' NT	Jo	First SEQ	-	Start Signal NO:	Pep	485		941		164		1508		206		229		133			195		128			253	
		5° NT		Start	Codon	485												133			195					253	
	5' NT 3' NT	Jo	Total Clone Clone	Seq.		655		1286		626		1765		721		1024		1210			1428		1633			1406	
	5' NT	Jo	Clone	Seq.		I		735		1		1391		1		1		1			1012		13			-	
				NT	Seq.	655		1286		979		2186		721		1024		1210			1445		1633			1406	
	N	SEQ		NO:	X	399		400		401		402		403		404		405			406		407			408	
					Vector	Uni-ZAP XR		Uni-ZAP XR			Uni-ZAP XR		Uni-ZAP XR			Uni-ZAP XR											
		ATCC	Deposit	No:Z	and Date	203959	04/26/99	203959	04/26/99	203918	04/08/66	203959	04/26/99	203959	04/26/99	203918	04/08/99	PTA-	181	66/L0/90	203959	04/26/99	PTA-	791	09/27/99		04/08/99
				cDNA	Clone ID	HSLHX15		HSNAP85		HSNAZ09		HSNBM34		HSOAH16		HSQBF66		HSQDO85			HSQES57		HSRBE06			HSSDI26	
				Gene	No.	389		390		391		392		393		394		395			396		397			398	

		Last	AA	of	ORF	62		!	09		125		38		59		14			33		27		4		137		
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		First	AA of	Secreted	Portion	11			97		20		81		38					22		20				11		
	Last	AA	Jo	Sig	Pep	16			25		19		17		34					21		19				16		
·	AA First Last	SEQ AA		Sig	Pep	_					1		-		Т		1			-1				1		I		
		SEQ	А	N	Y	913			914		915		916		917		918			919		920		921		922		
5' NT	of	First	AA of	Signal NO:	Pep	28			184		264		245		380		211			232		366		306		39		
		5' NT	of	Start	Codon	28							245		380					232		366						
	3, NT	of	Clone Clone	Seq.		1274			1053		1133		1954		710	·	2206			956		1198		2174		1764		
	5' NT 3' NT	of	Clone	Seq.		1			1		85		1		250		1			1		-		1		1		
			Total	N	Seq.	1282			1053		1238		1954	,	874		2206			926		1198		2174		1764		
	NT	SEQ	Ω	NO:	X	409			410		411		412		413		414			415		416		417		418		
					Vector	Uni-ZAP XR		•	Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		pCMVSport	3.0	Uni-ZAP XR			pCMVSport	3.0	Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		
		ATCC	Deposit	No:Z	and Date	PTA-	181	66/L0/90	203959	04/26/99	203959	04/26/99	203918	04/08/69	203959	04/26/99	PTA-	795	09/27/60	203918	04/08/99	203959	04/26/99	203959	04/26/99	PTA-	181	66/20/90
				cDNA	Clone ID	HSSEA64			HSSEF77		HSSFE38		HSSGJ58		HSWBE76		HSXCP38			HSYBI06		HT1SC27		HT3BF49		HT4FV41		
				Gene	No.	399			400		401		402		403		404			405		406		407		408		

		Last	AA	of	ORF	20	31		4		20	T	22	_	99		23			2		45	,	36		32	
		يًا		_	$\neg$	Ň	6		7		2	4	7	_	Ň		2			3		4		3		3	
		First	AA of	Secreted	Portion	18	24						21		23	!	07					61		25		20	
	Last	AA	of	Sig	Pep	17	23						20		22		61					18		24		61	
	First	AA	of	Sig	Pep	1	1		1		1		-		_		Ţ	•		1		1		1		1	
	AA	SEQ	Д		Y	923	924		925		926		927		928		926			930		931		932		933	
5' NT	of	First	AA of	Signal	Pep	228	135		632		151		1017		196		211			325		287		260		261	
		5' NT	of	Start	Codon		135		632		151		1017		196		211			325				260		261	
	3, NT	Jo	Clone Clone	Seq.		682	1743		1623		825		2221		1662		2055			678		1247		1587		2179	
	5' NT 3' NT	of	Clone	Seq.		59			-		1		27		106		1			1		1		1		1	
			Total	Ŋ	Seq.	682	1743		1623		825		2221		1662		2055			829		1247		1587		2179	
	N	SEQ	А	NO:	X	419	420		421		422		423		424		425			426		427		428		429	
					Vector	Uni-ZAP XR	Uni-ZAP XR	,	Uni-ZAP XR		pSport1		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR			Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR	
		ATCC	Deposit	No:Z	and Date	203959	203959	04/26/99	203918	04/08/69	203918	04/08/99	203959	04/26/99	203959	04/26/99	PTA-	181	66/L0/90	203959	04/26/99	203959	04/26/99	203918	04/08/99	203959	04/26/99
				cDNA	Clone ID	HTSFX79	HT5GR59		HTAEI78		HTDAA78	- 1	HTEAG62		HTECB02		HTECC15			HTEDF18		HTEDJ28		HTEDS12		HTEED26	
				Gene	No.	409	410		411		412		413		414		415			416		417		418		419	

		Last	AA	Jo	ORF	32	Т		$\neg$	_		323		37	Т	_		П	88		34		23		38		
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		First	AA of	Secreted	Portion	20						31		21					9		70				25		
	Last	AA	of	Sig	Pep	19						30		20					2		119				24		
	AA First Last	AA	of	Sig	Pep	_		1				1		-							_		1		-		
	AA	SEQ	А	NO:	Y	934		935		936		937		938		939			940		941		942		943		
5' NT	Jo	First SEQ	AA of	Signal NO:	Pep	259		262		262		182		493		173			280		170		101		171		
		5, NT	of	Start	Codon	259		262		262		182									170		101		171		
	3, NT	Jo	Clone	Seq.		5129		984		984		1263		908		981			1400		1504		1324		2116		
	5' NT 3' NT	Jo .	Clone Clone	Seq.		1		45		45		110		1					529		_		_		1		
			Total	N	Seq.	2167		1015		1273		1282		908		981			1402		1523		1324		2116		
	N	SEQ	Ω	NO:	X	430		431		432		433		434		435			436		437		438		439		
					Vector	Uni-ZAP XR		Uni-ZAP XR			Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR										
		ATCC	Deposit	No:Z	and Date	203959	04/26/99	203959	04/26/99	203959	04/26/99	203959	04/26/99	203959	04/26/99	PTA-	181	66/L0/90	203959	04/26/99	203959	04/26/99	203959	04/26/99	PTA-	181	66/L0/90
				cDNA	$\overline{}$	HTEED26		HTEEF26		HTEEF26		HTEEW69		HTEGS07		HTEGS11			HTEHA56		HTEHU59		HTEJD29		HTEKM46		
				Gene	No.	420		421		422		423		424		425			426		427		428		429		

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		ATCC		SEQ (		o i			FIISL	) (	AA J	74 70	Je II.I	- ast
		Deposit				Clone Clone	Clone		AA OI	<b>∃</b>			AA OI	AA
Gene	cDNA	No:Z		Ö	Ę	Seq.	Sed.		_	_	S1g		_	o I
Š.	Clone ID	and Date	Vector	X	Seq.			Codon	Pep	>	Pep	Pep	Portion	SF
430	HTEMQ17	203959	Uni-ZAP XR	440	1768		1768	446	446	944				12
		04/26/99												1
431	HTENR63	PTA-	Uni-ZAP XR	441	1591	-	1591	132	132	945		70	21	99
		792												
		09/27/99												
432	HTGGM44	203959	Uni-ZAP XR	442	3016	-	2761	179	179	946		18	19	<b>2</b>
		04/26/99												
433	HTHBZ06	203959	Uni-ZAP XR	443	623	193	619	318	318	947				<del>-</del>
		04/26/99												
434	HTLAP64	203918	Uni-ZAP XR	444	1092	-	1092	173	173	948		19	70	70
		04/08/99												
435	HTLBT80	203959	Uni-ZAP XR	445	2101	817	1881	912	912	949	_	27		129
		04/26/99												,
436	HTLDA84	203918	Uni-ZAP XR	446	1444	-	1444		225	950	-			13
		04/08/99												
437	HTLDN29	203959	Uni-ZAP XR	447	1374	-	1348	175	175	951	_	23	74	33
		04/26/99												
438	HTLDU78	203918	Uni-ZAP XR	448	1318	-	1318	219	219	952				×
		04/08/99												
439	HTLEC82	203959	Uni-ZAP XR	449	1260	217	1119	530	230	953	_	34	35	36
		04/26/99												,
440	HTLEM16	203959	Uni-ZAP XR	450	1915	1158	1755	1220	1220	954		27	78	69
		04/70/99												

		Last	AA	of	ORF	207		6		31	T	23		37		38		75		75		75		75		75	
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		First	AA of	Secreted	Portion	31				6				17		34		20		20		70		20		20	
	Last	AA	of	Sig	Pep	30				∞				16		33		19		19		19		19		19	
	First Last	AA	of	Sig	Pep			-		_		_		_		-				_		_				_	
	AA	SEQ	白	NO:	Υ	955		1018		926		957		958		959		096		961		962		696		964	
5' NT	Jo	First	AA of	Signal	Pep	205		91		209		340		1802		933		642		644		644		644		644	
		5' NT	Jo	Start	Codon	205		91				340		1802		933		642		644		644		644		644	
	3, NT	Jo	Clone	Seq.		1070		1065		1160		1159		2377		1968		1100		1033		1033		1033		1033	
	5' NT 3' NT	of	Clone Clone	Seq.		1		1				1		1205		098		140		142		142		142		142	
			Total	K	Seq.	1070		1065		1160		1159		2377		1968		1100		1081		1044		1081		1081	
	Z	SEQ	А	SO.	X	451		514		452		453		454		455		456		457		458		459		460	
					Vector	Uni-ZAP XR		Uni-ZAP XR	,	Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR	
		ATCC	Deposit	No:Z	and Date	203918	04/08/99	203918	04/08/66	203918	04/08/99	203979	04/29/99	203959	04/26/99	203959	04/26/99	203959	04/26/99	203959	04/26/99	203959	04/26/99	203959	04/26/99	203959	04/26/99
				cDNA	Clone ID	HTLEV48		HTLEV48		HTLFA13		HTLFI73		HTLGI89		HTLIF11		HTLIF12									
				Gene	No.	441		441		442		443		444		445		446		447		448		449		450	

		Last	A	Jo	ORF	75		30		21		27		34		71		32		118			42		— 63	
		<u>~</u>	AA		$\overline{}$	7		n		_	_	-		$\frac{\omega}{\omega}$	_	_		$\omega$	_	_			4	$\dashv$	<u> </u>	
		First	AA of	Secreted	Portion	20		16		17		18		25		19		22		40			24		9	
	First Last	AA	oę	Sig	Pep	19		15		16		17		24		18		21		36			23		2	
	First	AA	of	Sig	Pep	1		_		-		_		—				<del></del>					1		-	
	AA	SEQ	А	NO:	Y	965		996		296		896		696		970		971		972			813		974	
5' NT	Jo	First	AA of	Signal	Pep	644		193		534		61		68		228		103		201			314		439	
		5' NT	of	Start	Codon	644				534		61		68				103		201				_		
	3, NT	of	Clone	Seq.		1033		1006		1148		1258		1200		1652		1981		1640			9//		727	
	5' NT 3' NT	Jo	Clone Clone	Seq.	_	142		_		295		1		1		1		_		-			138		1	
			Total	K	Seq.	1081		1006		1160		1258		1200		1652		1981		1640			9/1		727	
	Z	SEQ	А	NO:	X	461		462		463		464		465		466		467		468			469		470	
					Vector	Uni-ZAP XR		pBluescript	SK-	pBluescript	SK-	Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR		Uni-ZAP XR			Uni-ZAP XR		Uni-ZAP XR	
		ATCC	Deposit	No:Z	and Date	203959	04/26/99	203918	04/08/69	203959	04/26/99	203959	04/26/99	203918	04/08/99	203918	04/08/66	203918	04/08/69	PTA-	181	66/L0/90	203959	04/26/99	203918	04/08/60
				cDNA	$\overline{}$	HTLIF12		HTNAM63		HTNBK13		HTOAI50		HTOAM11		HTODH57		<b>Е8НДОТН</b>		HTOEV16			HTOGR38		HTOH021	
				Gene	No.	451		452		453		454		455		456		457		458			459		460	

ATCC SEQ
Deposit ID
NO:
and Date   Vector   X   Seq.
PTA-   Uni-ZAP XR   471   1860
181
203959 Uni-ZAP XR 472 1854
04/26/99
203959 Uni-ZAP XR 473 1947
04/26/99
203959 Uni-ZAP XR   474   2078   04/26/99
203918 pBluescript 475 1257
04/08/99
PTA- Uni-ZAP XR   476   1504   181
66/10/90
203959 Uni-ZAP XR 477 1973
04/26/99
203918 Uni-ZAP XR 478 1880 04/08/99
203918 pSport1 479 1361
•
203959 Uni-ZAP XR 480 1921
104/20/22

								<del>,                                    </del>				
Last	AA	ORF	22	18	151	132	99	59	29	16	78	37
First	AA of Secreted	Portion	11		19	18	25	23	18		56	23
Last AA	of Sio	от Рер	10		18	17	24	22	17		28	22
First Last AA AA	of Sio	от Рер	1	-	1	1	-	1	1	_	1	-
AA SEQ	Βġ	Y	586	986	<i>L</i> 86	886	686	066	991	992	993	994
5' NT of First	AA of Signal	Dep	229	359	<b>59</b>	49	216	178	192	108	330	319
5' NT	of	ı		359	99			178	192	108	330	319
3' NT of	Clone	ocq.	1135	908	1400	1140	1162	686	1861	1187	875	1640
5' NT 3' NT of		ocy.	1	106	159	_	-	-	_	12	62	189
	Total	Seq.	1211	820	1441	1140	1162	686	1861	1187	884	1652
NT SEQ	Αġ	X	481	482	483	484	485	486	487	488	489	490
		Vector	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR	Uni-ZAP XR
ATCC	Deposit	and Date	PTA- 181 06/07/99	203959	203979 04/29/99	PTA- 181 06/07/99	203959	203918 04/08/99	PTA- 181 06/07/99	203959	203959	203959 04/26/99
	V NO.	Clone ID	HTXDB22	HTXDC38	HTXDC77	HTXDD61	HTXDG92	HTXET11	HTXFA72	HTXJY08	HTXKF95	HTXMZ07
	, and a	No.	471	472	473	474	475	476	477	478	479	480

ATCC  ODNA  No.Z  Clone ID  OH/26/99  HUKBT67  203959  HUKDY82  Clone ID  OH/26/99  HUKDY82  Clone ID  OH/26/99  HUKDY82  Clone ID  OH/26/99  HUKDY82  OH/26/99  HUKDY82  OH/26/99  II  HUKDY82  OH/26/99  OH/26/99  HUSGU40  OH/26/99  PSport1  OH/26/99  HUSGU40  OH/26/99  HUMAAII2  OH/26/99  HWAAII2  OH/26/99  OH/26/99  HWABQ70  OH/26/99  OH/26/99  OH/26/99  OH/26/99  OH/26/99  HWABQ70  OH/26/99  OH										Z, NT					
ATCC Deposit Deposit Deposit Deposit Deposit Deposit No:Z					Z		S, NT	3, NT		of	AA	First Last	Last		
CDNA         No.Z         NO.Z         NO.Z         NO.Z         NO.Z         NO.Z         NO.Z         Seq.         Seq.         Seq.         Start         Signal         Codon         Pep           HUFCL31         203959         pSport1         491         1460         1         1460         287         287           HUKBTG         203959         Lambda ZAP         492         2069         74         2052         273           HUKDF20         203918         Lambda ZAP         493         1105         1         1105         214         214           HUKDY82         203918         Lambda ZAP         494         1435         1         1435         187         187           HUKDY82         203918         Lambda ZAP         494         1435         1         1435         187         187           HUKDY82         203918         Lambda ZAP         495         3342         1         3342         74         74           HUSCI14         PSport1         496         1008         65         1008         350         350           HUSGU40         203959         pSport1         497         1054         1         1827         196			ATCC		SEQ		of		5' NT	First	SEQ	AA	AA	First	Last
cDNA         No:Z         NO:Z         NO:Z         Seq.         Seq.         Start         Signal Colone           Clone ID         and Date         Vector         X         Seq.         Codon         Pep           HUFCL31         203959         pSport1         491         1460         1         1460         287           HUKBT67         203959         Lambda ZAP         492         2069         74         2052         273           HUKDF20         203918         Lambda ZAP         493         1105         1         1105         214         214           HUKDPS0         203918         Lambda ZAP         494         1435         1         1435         187         187           HUKDYSU14         PTA-         Lambda ZAP         495         3342         1         3342         74         74           HUSCJ14         PTA-         Lambda ZAP         495         3342         1         3342         74         74           HUSCJ14         PTA-         Lambda ZAP         495         1054         1         1054         200           HUSCJ14         PTA-         Lambda ZAP         495         1054         1         1054 <t< td=""><td></td><td></td><td>Deposit</td><td></td><td>А</td><td>Total</td><td>Clone</td><td>Clone</td><td>of</td><td>AA of</td><td>А</td><td></td><td></td><td>AA of</td><td>AA</td></t<>			Deposit		А	Total	Clone	Clone	of	AA of	А			AA of	AA
Clone ID         and Date         Vector         X         Seq.         Codon         Pep           HUFCL31         203959         pSport1         491         1460         1         1460         287           HUKBT67         203959         Lambda ZAP         492         2069         74         2052         273           HUKDF20         203918         Lambda ZAP         493         1105         1         1105         214         214           HUKDF82         203918         Lambda ZAP         494         1435         1         1435         187         187           HUKDP82         203918         Lambda ZAP         494         1435         1         1435         187         187           HUKDP82         203918         Lambda ZAP         495         3342         1         143         144           HUSCI14         PTA-         Lambda ZAP         495         3342         1         144         74           HUSGL67         203918         pSport1         496         1068         65         1068         350           HUSGU40         203959         pSport1         496         165         1         1876         196      <	ene	cDNA	No:Z		SON:	Z	_	Seq.		Signal	$\sim$	Sig			ot
HUFCL31         203959         pSport1         491         1460         1         1460         287           HUKBT67         203959         Lambda ZAP         492         2069         74         2052         273           HUKDF20         203918         Lambda ZAP         494         1435         1         1455         187         187           HUKDP82         203918         Lambda ZAP         494         1435         1         1435         187         187           HUKDP82         203918         Lambda ZAP         495         3342         1         1435         187         187           HUSCJ14         PTA-         Lambda ZAP         495         3342         1         1435         187         144           HUSCJ14         PTA-         Lambda ZAP         495         3342         1         3342         74         74           HUSGL67         203918         pSport1         496         1008         65         1008         350         350           HUVBIRS         203959         pSport1         496         1827         1         1827         196         196           HUVBBQ70         203959         pCMVSport         50<	j.		and Date		X	Seq.			Codon	Pep	X	Pep	Pep	Portion	ORF
HUKBT67 203959 Lambda ZAP 492 2069 74 2052 273  HUKDF20 203918 Lambda ZAP 493 1105 1 1105 214 214  HUKDY82 203918 Lambda ZAP 494 1435 1 1435 187 187  HUSCI14 PTA- Lambda ZAP 495 3342 1 3342 74 74  HUSCL67 203918 pSport1 496 1008 65 1008 350 350  HUSGU40 203959 pSport1 497 1054 1 1054 500  HUSR18 203959 pSport1 498 876 1 876 83 83  HUVDJ48 203918 Uni-ZAP XR 499 1827 1 1837 196 196  HWAAI12 203959 pCMVSport 500 3303 1 1838 223 223  HWBBQ70 203959 pCMVSport 501 1948 1 1948 222 222	481		203959		491	1460	1	1460		287	995	_			76
HUKDF20         203918         Lambda ZAP         493         1105         1         1105         214         214           HUKDY82         203918         Lambda ZAP         494         1435         1         1435         187         187           HUKDY82         203918         Lambda ZAP         495         3342         1         3342         74         74           HUSCJ14         PTA-         Lambda ZAP         495         3342         1         3342         74         74           HUSGL67         203918         pSport1         496         1008         65         1008         350         350           HUSGU40         203959         pSport1         497         1054         1         876         83         83           HUVDJ48         203959         pSport1         498         876         1         876         83         83           HWAAII2         203918         Uni-ZAP XR         499         1827         1         196         196           HWABQ70         203959         pCMVSport         501         1948         1         1948         222         222	182	HUKBT67	203959		492	2069	74	2052		273	966	-	21	22	39
HUKDF20         203918         Lambda ZAP         493         1105         1         1105         214         214           04/08/99         II         II         1435         1         1435         187         187           HUKDY82         203918         Lambda ZAP         495         3342         1         1435         187         187           HUSCJ14         PTA-         Lambda ZAP         495         3342         1         3342         74         74           HUSCJ14         PTA-         Lambda ZAP         495         3342         1         3342         74         74           HUSCJ16         PSport1         496         1008         65         1008         350         350           HUSGL67         203959         pSport1         497         1054         1         876         83         83           HUVDJ48         203918         Uni-ZAP XR         499         1827         1         1827         196         196           HWAAI12         203959         pCMVSport         500         3303         1         1838         223         223           HWBBQ70         203959         pCMVSport         501         1			04/26/99	П											
HUKDY82       Lambda ZAP       494       1435       1       1435       187       187         HUSCJ14       PTA-       Lambda ZAP       495       3342       1       3342       74       74         HUSCJ14       PTA-       Lambda ZAP       495       3342       1       3342       74       74         HUSGL67       203918       pSport1       496       1008       65       1008       350       350         HUSGL67       203959       pSport1       497       1054       1       876       83       83         HUSGU40       203959       pSport1       498       876       1       876       83       83         HUVDJ48       203918       Uni-ZAP XR       499       1827       1       1827       196       196         HWAAI12       203959       pCMVSport       500       3303       1       1838       223       223         HWBBQ70       203959       pCMVSport       501       1948       1       1948       1       1948       222       222	183	HUKDF20	203918	Lambda ZAP	493	1105		1105	214	214	266	_	70	21	33
HUKDY82         203918         Lambda ZAP         494         1435         1         1435         187         187           HUSCI14         PTA-         Lambda ZAP         495         3342         1         3342         74         74           1838         II         R         1838         II         8         74         74           HUSGU67         203918         pSport1         496         1008         65         1008         350         350           HUSGU40         203959         pSport1         497         1054         1         876         83         83           HUVDJ48         203959         pSport1         498         876         1         876         83         83           HUVDJ48         203918         Uni-ZAP XR         499         1827         1         1827         196         196           HWAAI12         203959         pCMVSport         500         3303         1         1838         223         223           HWBBQ70         203959         pCMVSport         501         1948         1         1948         222         222			04/08/99	П											
HUSCJ14         PTA-         Lambda ZAP         495         3342         1         3342         74         74           HUSCJ14         PTA-         Lambda ZAP         495         3342         1         3342         74         74           605/09/00         05/09/00         PSport1         496         1008         65         1008         350         350           HUSGL67         203959         PSport1         497         1054         1         1054         500           HUSIR18         203959         PSport1         498         876         1         876         83         83           HUVDI48         203959         PCMVSport         499         1827         1         1827         196         196           HWAAI12         203959         PCMVSport         500         3303         1         1838         223         223           HWBBQ70         203959         PCMVSport         501         1948         1         1948         222         222	184	HUKDY82	203918			1435	1	1435	187	187	866	<del>-</del>	17	18	32
HUSCJ14         PTA-         Lambda ZAP         495         3342         1         3342         74         74           1838         II         1838         II         65/09/00         65/1008         65         1008         350         350           HUSGL67         203959         pSport1         496         1008         65         1008         350         350           HUSGU40         203959         pSport1         497         1054         1         1054         500           HUSIR18         203959         pSport1         498         876         1         876         83         83           HUVDJ48         203918         Uni-ZAP XR         499         1827         1         186         196           HWAAI12         203959         pCMVSport         500         3303         1         1838         223         223           HWBBQ70         203959         pCMVSport         501         1948         1         1948         222         222			04/08/99	П											
HUSGL67       203918       pSport1       496       1008       65       1008       350       350         HUSGU40       203959       pSport1       497       1054       1       1054       500         HUSGU40       203959       pSport1       498       876       1       876       83       83         HUSIR18       203959       pSport1       498       876       1       876       83       83         HUVDJ48       203918       Uni-ZAP XR       499       1827       1       1827       196       196         HWAAI12       203959       pCMVSport       500       3303       1       1838       223       223         HWBBQ70       203959       pCMVSport       501       1948       1       1948       222       222	185	HUSCJ14	PTA-	Lambda ZAP	495	3342		3342	74	74	666	_	30	31	196
HUSGL67         203918         pSport1         496         1008         65         1008         350         350           HUSGU40         203959         pSport1         497         1054         1         1054         500           HUSGU40         203959         pSport1         498         876         1         876         83         83           HUSIR18         203959         pSport1         498         876         1         876         83         83           HUVDJ48         203918         Uni-ZAP XR         499         1827         1         1827         196         196           HWAAI12         203959         pCMVSport         500         3303         1         1838         223         223           HWBBQ70         203959         pCMVSport         501         1948         1         1948         222         222			1838	п											
203918       pSport1       496       1008       65       1008       350       350         203959       pSport1       497       1054       1       1054       500         203959       pSport1       498       876       1       876       83       83         203918       Uni-ZAP XR       499       1827       1       1827       196       196         203959       pCMVSport       500       3303       1       1838       223       223         203959       pCMVSport       501       1948       1       1948       1       1948       222       222			02/06/00												
HUSGU40       203959       pSport1       497       1054       1       1054       500         HUSIR18       203959       pSport1       498       876       1       876       83       83         HUVDJ48       203918       Uni-ZAP XR       499       1827       1       1827       196       196         HWAAI12       203959       pCMVSport       500       3303       1       1838       223       223         HWBBQ70       203959       pCMVSport       501       1948       1       1948       1       1948       222       222	981	L9TDSOH	203918	pSport1	496	1008	65	1008	350	350	1000	_	21	22	47
HUSGU40         203959         pSport1         497         1054         1         1054         500           HUSIR18         203959         pSport1         498         876         1         876         83         83           HUVDJ48         203918         Uni-ZAP XR         499         1827         1         1827         196         196           HWAAI12         203959         pCMVSport         500         3303         1         1838         223         223           HWBBQ70         203959         pCMVSport         501         1948         1         1948         222         222			04/08/66												
HUSIR18       203959       pSport1       498       876       1       876       83       83         HUVDJ48       203918       Uni-ZAP XR       499       1827       1       1827       196       196         HWAAI12       203959       pCMVSport       500       3303       1       1838       223       223         HWBBQ70       203959       pCMVSport       501       1948       1       1948       1       1948       222       222	187	HUSGU40	203959	pSport1	497	1054	1	1054		200	1001	-	70	21	46
HUSIR18         203959         pSport1         498         876         1         876         83         83           HUVDJ48         203918         Uni-ZAP XR         499         1827         1         1827         196         196           HWAAI12         203959         pCMVSport         500         3303         1         1838         223         223           HWBBQ70         203959         pCMVSport         501         1948         1         1948         222         222			04/26/99												
HUVDJ48       203918       Uni-ZAP XR       499       1827       1       1827       196       196         HWAAI12       203959       pCMVSport       500       3303       1       1838       223       223         HWBBQ70       203959       pCMVSport       501       1948       1       1948       222       222	88	HUSIR18	203959	pSport1	498	9/8	-	9/8	83	83	1002	_	16	17	22
HUVDJ48         203918         Uni-ZAP XR         499         1827         1         1827         196         196           04/08/99         190			04/26/99												
HWAAI12       203959       pCMVSport       500       3303       1       1838       223       223         HWBBQ70       203959       pCMVSport       501       1948       1       1948       222       222	681	HUVDJ48	203918	Uni-ZAP XR	499		1	1827	196	196	1003				S
HWAAI12         203959         pCMVSport         500         3303         1         1838         223         223           04/26/99         3.0         3.0         3.0         3.0         203959         pCMVSport         501         1948         1         1948         222         222			04/08/99												
04/26/99         3.0         3.0           HWBBQ70         203959         pCMVSport         501         1948         1         1948         222         222	96	HWAAI12	203959	pCMVSport	200	3303	_	1838	223	223	1004	-			53
HWBBQ70 203959 pCMVSport 501 1948 1 1948 222 222			04/26/99	3.0											
	161	HWBBQ70	203959		501		-	1948	222	222	1005	-	21	22	43
٠			04/26/99	3.0											

								5' NT					
		•	N		5' NT 3' NT	3, NT		Jo	AA	AA First Last	Last		
	ATCC		SEQ		of	Jo	5' NT	First	SEQ	AA	AA	First	Last
	Deposit		_	Total	Clone	Total Clone Clone	of	AA of ID	А	of	of	AA of	AA
cDNA	No:Z		NO:	N	Seq.	Seq.	Start	Start Signal NO:	SON:	Sig	Sig	Secreted	Jo
Clone ID	and Date	Vector	X	Seq.			Codon	Pep	Y	Pep	Pep	Portion	ORF
HWBCN36	203959	pCMVSport	505	1008	1	1008	378	378	1006		23	24	8
	04/26/99	3.0											
HWBDJ08	203959	pCMVSport	503	2085		2085	253	253	1007	_	53	30	20
	04/26/99	3.0											
HWBFX16	203959	pCMVSport	504	1497	1	1497		267	1008	-			m
	04/26/99	3.0											
HWDAC26	203959	pCMVSport	505	1958	1	1958	242	242	1009		25	56	35
	04/26/99	3.0											
HWDAG96	203959	pCMVSport	206	1147	300	1147	998	998	1010	_	18	19	32
	04/26/99	3.0											
HWDAJ01	203959	pCMVSport	202	781	1	781	288	288	1011	—			74
	04/26/99	3.0											
HWHPB78	203959	pCMVSport	508	1346		1346	700	200	1012		23	24	99
	04/26/99	3.0						- 1					
HYABC84	203959	pCMVSport	509	1338	89/	1238	1015	1015 1013	1013	-	28	29	62
	04/26/99	3.0											
HYABC84	203959	pCMVSport	510	1478	833	1306	1080	1080 1014	1014	-	78	29	62
	04/26/99	3.0											

# Deesooke oethor

#### TABLE 1B

					AA		Tissue Distribution		OMIM
Gene	Clone ID	Contig	SEQ ID	ORF	SEQ		Library code: count	Cytologic	Disease
No:		Ä	NO: X	(From-To)	D NO: Y	Predicted Epitopes	(see Table 4 for Library Codes)	Band	Reference(s):
	H6BSF56	762968	=	83 - 508	515	Asn-131 to Met-140.	AR089: 53, AR060: 29 L0599: 4, L0439: 3, L0777: 3, H0253: 2, H0520: 2, L0754: 2, L0745: 2, L0759: 2, H0556: 1, H0657: 1, S0116: 1, H0450: 1, S0418: 1, S0046: 1, S0222: 1, H0492: 1, S0049: 1, H0123: 1, H0050: 1, H0051: 1, H0615: 1, S0036: 1, H0494: 1, L0805: 1, L0776: 1, S0126: 1, H0435: 1, H0670: 1, S0028: 1, L0747: 1,		
2	Н6ЕDМ64	841331	12	1448 - 1468	516		AR060: 22, AR089: 16 H033: 6, H0556: 5, H0255: 5, H0547: 5, H0618: 4, H0581: 4, H0553: 4, H0135: 4, L0783: 4, S0358: 3, S0222: 3, H0617: 3, L0769: 3, H0521: 3, H0617: 3, L0769: 3, H0402: 2, H0619: 2, H0549: 2, H0592: 2, H0253: 2, S0474: 2, H0620: 2, H0181: 2, H0659: 2, H0181: 2, H0659: 2, H0561: 2, L0761: 2, L0764: 2, L0809: 2, H0520: 2, H0682: 2, S0330: 2, H0522: 2,		

L0751: 2, L0747: 2, L0750: 2, L0755: 2, L0596: 2, L0601: 2, H0624: 1, H0686: 1, H0295: 1, H0686: 1, H0483: 1, S0356: 1, S0342: 1, S0356: 1, S0342: 1, S0356: 1, H0483: 1, S0356: 1, H0483: 1, S0356: 1, H0483: 1, S0356: 1, H0486: 1, H0486: 1, H0486: 1, H0486: 1, H0486: 1, H0486: 1, H0687: 1, H0688: 1, H0413: 1, H0687: 1, L0796: 1, H0687: 1, L0796: 1, L0687: 1, L0689: 1, L0789: 1, L0789: 1, H0136: 1, S0011: 1, H0136: 1, S0196: 1 and	H0352: 1. L0809: 4, L0747: 4, L0794: 3, L0759: 3, S0046: 2, H0497: 2, H0559: 2, H0575:
	1 6
	517
	263 - 319
	13
	889401
	H6EEC72
	3

# osssose osaes

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·		
2, H0618: 2, H0050: 2, L0769: 2, L0766: 2, L0663: 2, H0521: 2, L0743: 2, L0748: 2, H0650: 1, H0650: 1, H0650: 1, H0650: 1, H0657: 1, H0650: 1, H0650: 1, H0650: 1, H0635: 1, H0635: 1, H0635: 1, H0640: 1, H0635: 1, H0069: 1, H0640: 1, L0770: 1, L0644: 1, L0770: 1, L0660: 1, L06	L0748: 4, H0457: 3 and S6022: 1.	AR251: 7, AR310: 6, AR265: 6, AR053: 6, AR060: 6, AR055: 5, AR312: 5, AR309: 5, AR273: 5, AR061: 5, AR206: 5, AR194: 5, AR186: 5, AR213: 4, AR052: 4, AR089: 4, AR253: 4, AR248: 4, AR253: 4, AR039: 3, AR243: 3, AR096: 3, AR039: 3, AR246: 3, AR104: 3, AR202: 3,
	Leu-6 to Ser-12.	Arg-14 to Ile-24.
	518	519
	135 - 371	250 - 327
	14	15
	584773	847112
	HACAB68	HACBJ56
	4	S

AR263: 3, AR204: 2, AR244: 1, AR249: 1 H0661: 1, S0045: 1, H0550: 1, S0280: 1, S0010: 1, H0028: 1, L0764: 1, L0803: 1, L0665: 1, S0053: 1, H0670: 1, L0748: 1, L0731: 1 and L0581: 1.	H0052: 6, S0002: 5, H0580: 3, S0051: 3, L0766: 3, L0439: 3, L0777: 3, L0361: 3, S0046: 2, H0619: 2, H0550: 2, S0280: 2, H0039: 2, S0142: 2, L0794: 2, L0775: 2, L0748: 2, L0754: 2, L0747: 2, L0758: 2, L0596: 2, H0170: 1, H0265: 1, H0566: 1, S0349: 1, H0661: 1, H0663: 1, S0420: 1, S0356: 1, S0354: 1, H0637: 1, S0222: 1, H0431: 1, H0586: 1, H0492: 1, H0641: 1, H0542: 1, H0253: 1, R0042: 1, H0053: 1, R0033: 1, H0213: 1, H0148: 1, H0561: 1, H0063: 1, S038: 1, T0042: 1, H0560: 1, H0561: 1, S0372: 1, S0450: 1, S0344: 1, S0426: 1, L0762: 1, L0770: 1, L0769: 1, L0662: 1, H0539: 1, H0521: 1, S0174: 1, L0742: 1, L0751: 1, L0779: 1, L0779: 1, L0779: 1, L0779: 1, L0779: 1, L0779: 1, L0779: 1, L0779: 1,
	Cys-2 to Leu-8.
	250 C
	217 - 342
	16
	847113
	HACBS22
	φ

# ogenous collect

1, H0668: 1 and H0506: 1.	AR089: 25, AR060: 15 L0759: 6, L0769: 5, H0052: 4, L0770: 4, L0809: 4, L0439: 4, L0752: 4, S0408: 3, L0751: 3, L0747: 3, L0779: 3, S0007: 2, H0351: 2, H0333: 2, H0427: 2, H0581: 2, L0662: 2, L0777: 2, H0543: 2, L0741: 2, L0777: 2, H0543: 1, H0729: 1, H0171: 1, H0254: 1, H0261: 1, H0592: 1, H0586: 1, H0261: 1, H0592: 1, H0586: 1, H0261: 1, H0592: 1, H0642: 1, H0261: 1, H0592: 1, H0642: 1, H0123: 1, S0140: 1, H0587: 1, H0266: 1, L0768: 1, L0772: 1, L0646: 1, L0761: 1, L0772: 1, L0646: 1, L0765: 1, L0772: 1, L0646: 1, L0765: 1, L0772: 1, L0749: 1, L0765: 1, L0773: 1, H0547: 1, S0126: 1, H0547: 1, S0126: 1, H0557: 1, L0743: 1, L0758: 1, L0743: 1, L0758: 1, L0778: 1,	H0427: 1	AR089: 12, AR060: 7 H0124: 28, H0013: 8, H0547: 4, H0144: 3, L0595:
	Pro-9 to Thr-14, Ser-37 to Trp-44, Gly-79 to Thr-85, Arg-88 to Lys-139. Signal of the service of		
	521	522	523
	250 - 666	347 - 439	238 - 300
	17	2	19
	839187	877773	847116
	HADDE71	נוחחחדו	HADMB15
	7	0	0

## osscos: coleol

3, H0390: 2, S0346: 2, H0012: 2, L0565: 2, L0777: 2, S0001: 1, S0282: 1, S0442: 1, H0619: 1, S0222: 1, H0333: 1, T0039: 1, H0546: 1, H0178: 1, H0566: 1, H0178: 1, H0292: 1, H0135: 1, H0591: 1, H0087: 1, H0100: 1, L0770: 1, L0531: 1, L0651: 1, L0543: 1, L0664: 1, H0520: 1, L0593: 1, L0599: 1 and H0352: 1, L0599: 1 and H0352: 1.	AR060: 7, AR089: 4 L0754: 4, L0777: 2, L0755: 2, S0010: 1, H0049: 1, L0163: 1, L0771: 1, L0775: 1 and L0776: 1.	AR089: 16, AR060: 11 S0010: 1 and H0616: 1.	AR089: 8, AR060: 6 L0766: 13, L0663: 5, L0439: 3, L0747: 3, L0750: 3, H0580: 2, H0486: 2, H0013: 2, S0250: 2, L0662: 2, L0768: 2, L0527: 2, L0647: 2, L0792: 2, L0779: 2, L0596: 2, L0592: 2, L0362: 2, H0543: 2, H0556: 1, S0114: 1, H0661: 1, H0402: 1, S0420: 1, H0676: 1, H0438: 1, H0600: 1, H0497: 1, S0010: 1, L0471:
·			
	524	525	526
	171 - 236	238 - 291	146 - 313
	20	21	22
	722205	637489	823543
	HAGBQ12	HAGDW20	HAGEG10
	10	=	12

1, H0083: 1, H0267: 1, H0316: 1, H0090: 1, H0591: 1, H0038: 1, H0040: 1, L0060: 1, L0667: 1, L0373: 1, L0803: 1, L0650: 1, L0774: 1, L0775: 1, L0555: 1, L0659: 1, L0791: 1, L0666: 1, L0664: 1, L0665: 1, H0520: 1, H0547: 1, H0684: 1, H0521: 1, H0436: 1, H0540: 1, L0740: 1, L0756: 1, L0755: 1, L0758: 1, H0445: 1, H0542: 1 and H0443: 1.	AR089: 15, AR060: 14 H0585: 12, L0439: 8, H0052: 7, H0251: 7, L0805: 7, L0776: 6, S0010: 5, L0803: 5, L0745: 5, L0809: 4, L0438: 4, L0779: 4, L0747: 3, S0222: 2, H0438: 2, T0010: 2, S6028: 2, L0455: 2, L0794: 2, L0790: 2, S0028: 2, L0794: 2, L0592: 2, H0650: 1, S0001: 1, S0420: 1, S0436: 2, L0592: 1, H0156: 1, T0082: 1, S0490: 1, H0263: 1, H0178: 1, H0156: 1, H0051: 1, S0049: 1, H038: 1, H0040: 1, S0386: 1, S0039: 1, L0351: 1, L0370: 1, L0770: 1, L0766: 1, L0774: 1, L0783: 1, L0788: 1, L0770: 1, L0665: 1, L0352:
	- HC14101010010141111
	527
	515 - 550
	23
	HAGEQ79 828055
	13

1, S0380: 1, L0740: 1, L0777: 1, L0755: 1 and L0759: 1.	AR060: 5, AR089: 3 L0438: 7, L0439: 6, L0747. 4, L0005: 3, S0360: 3, H0547: 3, S0222: 2, L0105: 2, S0002: 2, S0426: 2, L0794: 2, L0659: 2, L0664: 2, L0754: 2, L0758: 2, H0566: 2, H0170: 1, H0171: 1, H0656: 1, S0212: 1, H0580: 1, H0455: 1, H0069: 1, S0003: 1, H0039: 1, S0036: 1, L0471: 1, H0699: 1, S0422: 1, L0763: 1, L0638: 1, L0649: 1, L0646: 1, L0773: 1, L0646: 1, L0773: 1, L0646: 1, L0773: 1, L0646: 1, L0775: 1, L0647: 1, S0052: 1, H0144: 1, H0682: 1, H0555: 1, L0742: 1, L0750: 1, H0555: 1, L0742: 1, S0434: 1 and S0452: 1,	AR060: 9, AR089: 7 H0521: 5, L0777: 5, S0376: 4, H0156: 3, H0519: 3, H0436: 3, L0731: 3, H0656: 2, H0580: 2, H0036: 2, L0471: 2, H0090: 2, H0040: 2, H0551: 2, H0494: 2, S0438: 2, H0529: 2, L0809: 2, H0144: 2, S0374: 2,
	Met-1 to Lys-6.	
	528	529
	241 - 405	900 - 932
	42	25
	847120	773286
	HAGFS57	HAGHN57
	4	15

H0593: 2, H0170: 1, H0583: 1, H0650: 1, S0418: 1, S0358: 1, S0045: 1, H0619: 1, H0532: 1, H0643: 1, H0532: 1, H0643: 1, H0590: 1, S0346: 1, H0591: 1, H0590: 1, S0010: 1, S0346: 1, H0581: 1, H0231: 1, H0046: 1, H0123: 1, S0028: 1, H0687: 1, S0003: 1, S0214: 1, H0252: 1, H0615: 1, H0615: 1, H0645: 1, S0366: 1, H0634: 1, H0646: 1, S0464: 1, H0646: 1, S0426: 1, H0626: 1, S0426: 1, H0639: 1, H0520: 1, H0435: 1, S0328: 1, H0539: 1, H0539: 1, H0539: 1, H0704: 1, S0328: 1, H0539: 1, H0704: 1, S0027: 1, L0439:	1, L0750: 1, L0756: 1, L0757: 1, L0581: 1, L0595: 1, H0543: 1 and H0423: 1. AR194: 23, AR205: 21, AR206: 20, AR039: 18, AR246: 17, AR244: 14, AR243: 14, AR052: 13, AR245: 13, AR198: 12, AR310: 12, AR271: 12, AR310: 12, AR271: 10, AR053: 10, AR251: 10, AR273: 10, AR251: 10, AR273: 10, AR251: 10, AR309: 9, AR060: 9, AR309: 9, AR061: 7, AR249: 6, AR096: 6,
	230
	196 - 237
	26
	847013
	HAHEA15
	16

# DOSTOSE DOSTOS

L0756: 3, H0599: 2, L0750: 2, L0753: 2, L0775: 1, L0754: 1, L0755: 1 and L0759: 1.	H0560: 1, H0561: 1 and H0542: 1.	AR060: 184, AR089: 98 H0561: 1 and L0758: 1.	AR089: 8, AR248: 7, AR269: 6, AR299: 7, AR265: 6, AR249: 6, AR253: 6, AR202: 6, AR312: 6, AR060: 5, AR194: 5, AR060: 5, AR194: 5, AR0513: 4, AR053: 3, AR310: 3, AR246: 3, AR205: 3, AR261: 3, AR206: 3, AR271: 3, AR204: 2, AR271: 3, AR204: 2, AR055: 1, AR186: 1, AR057: 2, AR204: 4, H0040: 4, L0659: 4, H0171: 3, H0040: 4, H0521: 4, H0171: 3, H0040: 3, H0553: 3, H0543: 3, S0418: 2, S0360: 2, S0222: 2, H0013: 2, H0551: 2, H0574: 2, H0615: 2, H0617: 2, H0623: 2, L0660: 2, L0660	S0374: 2, S0380: 2, S0146:
1, 10, 10, 10, 10, 10, 10, 10, 10, 10, 1	, H	·	25, 86.	SC
	Leu-33 to Asp-38	Lys-89 to Glu-94.	Arg-24 to Trp-44, Leu-87 to Ser-93, Arg-119 to Trp-125, Pro-206 to Lys-211, Glu-280 to Trp-286.	
,	531	532	533	
	192 - 308	12 - 296	605 - 1684	
	27	28	29	
	534670	845601	866415	
	HAJAA47	HAJAY92	HAJBV67	
	17	18		

2, L0740: 2, L0731: 2, L0759: 2, S0436: 2, L0362: 2, H0556: 1, S0114: 1, T0049: 1, L0002: 1, S0282: 1, S0356: 1, S0354: 1, S0408: 1, S0410: 1, H0637: 1, H0722: 1, S0046: 1, H0722: 1, S0046: 1, H0731: 1, H0731: 1, H0731: 1, H0731: 1, H0736: 1, L0717: 1, H056: 1, L0471: 1, H0756: 1, L0471: 1, H0756: 1, L0471: 1, H0756: 1, S0003: 1, S0214: 1, H0688: 1, H0251: 1, H0559: 1, L065: 1, H075: 1, L0662: 1, L0766: 1, L0388: 1, L0774: 1, L075: 1, L0665: 1, H0519: 1, S0126: 1, H0683: 1, L0899: 1, L0665: 1, H0519: 1, S0126: 1, H0683: 1, L0899: 1, L0665: 1, H0519: 1, S0126: 1, H0652: 1, L0809: 1, L0665: 1, H0519: 1, S0126: 1, H0651: 1, L0899: 1, L0659: 1, L075: 1, H0595: 1, S0394: 1, L0586: 1, L0589: 1, L075: 1, S0194: 1,	H0425: 1, H0422: 1, S0284: 1, H0506: 1 and H0352: 1.	- 1	AR206: 3, AR263: 3, AR207: 3, AR312: 2,
			Asp-26 to Leu-32, Trp-62 to Asp-72,
•		534	535
		284 - 400	8 - 3511
		30	31
		827275	852204
		HAJCH70	HAOAG15
		20	21

### Descours.cereor

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AR053: 2, AR254: 2, AR205: 1, AR089: 1, AR033: 1, AR096: 1 L0759: 3, S0314: 2, L0744: 2, L0756: 2, L0755: 2, S0046: 1, H0391: 1, H0052: 1, H0050: 1, S0318: 1, S0338: 1, S0312: 1, L0766: 1 and H0144: 1.	AR089: 43, AR060: 31 H0617: 5, H0606: 2, L0744: 2, L0779: 2, H0295: 1, H0100: 1, S0440: 1, H0026: 1, L0762: 1, L0504: 1, L0769: 1, L0764: 1, L0662: 1, L0649: 1, L0804: 1, L0787: 1, L0666: 1, L0663: 1, H0520: 1, L0748: 1, L0751: 1, L0752: 1 and S0436: 1.	AR060: 7, AR089: 3 H0295: 5	AR060: 5, AR089: 3 L0758: 9, L0769: 4, H0556: 3, L0756: 3, H0486: 2, H0156: 2, H0040: 2, H0529: 2, L0766: 2, L0803: 2, L0659: 2, L0809: 2, L0565: 2, H0539: 2, L0748: 2, L0754: 2, L0777: 2, H0595: 2, L0595: 2, L0361: 2, S0114: 1, H0402: 1, S0358: 1, H0580: 1, S0222: 1, H0587: 1, H0497: 1, H0013: 1, H0427: 1, H0581: 1, H0251: 1, H04046: 1, H0009: 1, H0320: 1, H0594: 1, H026: 1, H0031: 1,
Gln-95 to His-101, Thr-158 to Thr-164, Phe-222 to Glu-227, Asn-234 to Thr-245, Gly-256 to Glu-266, Gly-277 to Glu-283, Arg-310 to Ser-317, Ser-327 to Phe-333, Ser-360 to Ser-366.			Lys-42 to Asp-54.
	536	537	538
	250 - 321	262 - 273	18 - 224
	32	33	<del>2</del> 6
	688037	633730	839468
	HAQAI92	HAQCE11	HATBI94
	22	23	42

		:		
L0055: 1, H0376: 1, H0634: 1, S0038: 1, H0100: 1, L0667: 1, L0771: 1, L0804: 1, L0790: 1, L0791: 1, L0793: 1, L0665: 1, H0144: 1, H0519: 1, S0126: 1, H0682: 1, H0659: 1, H0521: 1, S0404: 1, L0759: 1, S0404: 1, L0759: 1, S0436: 1 and L0591: 1.	L0749: 3, H0156: 2, H0341: 1 and L0754: 1.	AR060: 3, AR089: 1 H0156: 1 and H0038: 1.	AR089: 17, AR060: 10 S6026: 1, H0156: 1 and S0426: 1.	AR060: 6, AR089: 4 L0439: 11, L0740: 11, H0046: 10, H0556: 8, H0052: 7, L0766: 7, S0222: 6, H0617: 6, S0049: 5, H0620: 5, H0144: 5, L0741: 5, L0747: 5, L0731: 5, S0278: 4, L0743: 4, L0743: 4, L0742: 4, L0743: 4, L0748: 4, H0657: 3, H0599: 3, H0618: 3, S0010: 3, H0650: 3, S0051: 3, S6028: 3, H0266: 3, H0551: 3, H0494: 3, S0144: 3, H0559: 3, L0751: 3, L0752: 3, L0663: 3, S0330: 3, L0751: 3, L0759: 3, H0665: 2, H0333: 2, H0457: 2, H0041: 2, T0010:
			Lys-8 to Trp-13.	Val-23 to Glu-28.
	539	540	541	542
	268 - 396	296 - 409	271 - 324	93 - 221
	35	36	37	38
	631172	826098	280805	836056
	HATCB45	HATCD80	HATC103	HATEH20
	25	26	27	78

2, S0003: 2, T0006: 2,	S0364: 2, H0124: 2, S0366:	2, H0135: 2, L0775: 2,	L0809: 2, L0789: 2, H0660:	2, L0753: 2, L0757: 2,	L0758: 2, L0485: 2, L0599:	2, L0601: 2, H0265: 1,	S0040: 1, H0650: 1, H0341:	1.50212.1.50282.1	0662, 1 H0629, 1 c0419.	HU003: 1, HU038: 1, 50418:	1, S0420: 1, S0360: 1,	S0408: 1, L0149: 1, H0208:	1, S0132: 1, H0370: 1,	L0623: 1. H0013: 1. H0427:	S0280: 1. H0156: 1	H0097 1 H0575 1 H0036	LIOSO0, 1 CO246, 1	1, H0390: 1, 30346: 1,	H0318: 1, H0196: 1, H0596:	1, H0597: 1, H0231: 1,	H0009: 1, N0006: 1, L0471:	1, H0012: 1, H0024: 1,	H0373: 1, H0051: 1, H0083:	1, H0292: 1, H0428: 1,	H0604: 1, H0553: 1, H0181:	1, H0169: 1, H0163: 1,	H0090: 1, T0067: 1, H0264:	1, S0038: 1, S0386: 1,	S0112: 1, L0564: 1, H0561:	, S0142: 1, S0344: 1,	L0770: 1, L0769: 1, L0637:	, L0761: 1, L0372: 1,	L0374: 1, L0521: 1, L0626:	1, L0533: 1, L0803: 1,	L0376: 1, L0806: 1, L0805:	, L0655: 1, L0783: 1,	L0529: 1, L0666: 1, S0374:	, H0520: 1, H0547: 1,	H0519: 1, H0658: 1, S0380:
2,	<u> </u>	2,	<u> </u>	2,	<u> </u>	2,	SC			<u>ii</u>	1,	)S	1,				<u> </u>		)H	1	)H	1,	)H	1,	)H	1,	)H	$\frac{1}{1}$	OS SO	$\frac{1}{1}$ ,		1	<u>  L0</u>		<u> </u>	1,1	T0	1,	HC
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# Doomoom, Doelnor

1, H0521: 1, H0522: 1, H0696: 1, H0436: 1, L0609: 1, L0744: 1, L0745: 1, L0749: 1, L0777: 1, H0444: 1, L0480: 1, L0584: 1, L0595: 1, S0011: 1, H0422: 1 and H0008: 1.	AR089: 1 L0809: 4, L0766: 3, L0439: 3, H0624: 2, H0411: 2, L0794: 2, L0756: 2, L0731: 2, L0005: 1, H0599: 1, L0471: 1, S0051: 1, T0010: 1, H0266: 1, S0150: 1, L0637: 1, L0765: 1, L0803: 1, L0783: 1, H0144: 1, H0672: 1, S0392: 1, L0748: 1, L0779: 1, L0777: 1 and L0759: 1.	ARÖ89: 8, ARO60: 4 H0013: 8, L0805: 5, H0716: 4, S0010: 4, H0052: 4, H0144: 4, H0615: 3, H0547: 3, L0747: 3, H0645: 2, S0049: 2, H0009: 2, L0769: 2, L0776: 2, L0665: 2, H0519: 2, H0658: 2, H0660: 2, L0602: 2, H0555: 2, L0439: 2, L0750: 2, L0597: 2, H0136: 2, H0423: 2, H0624: 1, H0171: 1, H0717: 1, S0402: 1, H0294: 1, S0114: 1, S0116: 1, H0341: 1, S0212: 1, H0483: 1, H0664: 1, S0360: 1, S0046: 1, H0619: 1, H0411: 1, H0369: 1, S0222: 1, H0338: 1, H0486: 1, H0156:
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	838799	1300785
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	561935	705047	589515
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	995	270	571
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	866160	778066	834801
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	Glu-59 to Gln-65, Lys-90 to Val-95, Glu-205 to Ser-211. Flu-205 to	
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		86
	828945	793774
	HCRBF72	HCRNF78
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		Gly-14 to Asp-21.	
		604	605
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	589520	637986	790277
	HCUAF85	HCUCF89	HCUCK44
	68	06	91

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S0436:	L0774:	3, L074	L0731:	2, H063	80408:	2, H005	H0416:	2. H04	1,0770	7 1 064	7, 200	9 8043	2, 5042	S0404:	2, L075	L0596:	1. H01	H0657:	1 - S044	\$0410:	1. H060	H0486	1, H05	6000H	1, S005	H0615	1, H00	H0087	1, 504	80210:	1. LO7	L0372:	1. L07	1.0662:	1, L06	L0550.	1, L06
			-								-																			11.00							
		-																																			

L0782: 1, L0787: 1, S0374: 1, H0659: 1, H0658: 1, S0378: 1, H0710: 1, S0152: 1, H0696: 1, H0704: 1, L074: 1, L0756: 1, L0779: 1, L0780: 1, L0759: 1, L0759: 1, L0761: 1, L0759: 1, L0601: 1, L0603: 1, S0196: 1 and H0352: 1.	H0052: 3, S3012: 2, L0754: 2, H0402: 1, H0413: 1, S0374: 1, L0438: 1, L0748: 1 and L0740: 1.	H0305: 1	AR089: 3, AR060: 1 H0305: 3, L0439: 3, L0740: 3, L0581: 2, H0589: 1, H0156: 1, S0346: 1, H0318: 1, S0049: 1, H0052: 1, L0157: 1, T0010: 1 and L0438: 1.	H0305: 9, H0589: 2 and S0001: 1.	AR089: 9, AR060: 8 H0616: 4, L0803: 3, H0555: 3, H0038: 2, L0809: 2, L0439: 2, L0759: 2, L0005: 1, S0049: 1, H0569: 1, S0050: 1, L0163: 1, S0003: 1, L0771: 1, L0649: 1, L0804: 1, L0774: 1, L0775: 1, L0784: 1, L0659: 1, L0788: 1, L0664: 1, L0438: 1, H0648: 1, S0330: 1, L0602: 1, L0744: 1, L0749: 1, L0745: 1, L0747:
	Met-1 to Ser-6, Gln-32 to Asn-39.				
	909	607	809	609	610
	256 - 402	410 - 427	282 - 350	333 - 368	48 - 128
	102	103	101	105	106
	835082	535893	651316	834722	695710
·	HCUDD64	HCWAE64	НСМЕЛ39	HCWUL09	НДНАА42
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	,	612
		172 - 339
	107	108
	553622	840358
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	Arg-63 to Phe-72, Ile-114 to Phe-120.	
	613	614
	23 - 385	279 - 341
	109	110
	897277	587265
	HDPDI72	HDPDJ58
	66	001

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		616
		220 - 378
		112
	853513	790189
	HDPFF10	HDPFU43
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	779450	801947
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	777493
	HDPOC24
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	189 - 377	109 - 159
	117	118
	745377	838594
	HDPOL37	HDP0076
	107	108

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			Glu-21 to Leu-26, Pro-34 to Ser-41.	
	624	625	979	627
	220 - 336	395 - 484	61 - 186	129 - 275
	120	121	122	123
	684292	778405	801896	852328
	HDPP030	HDPPW82	HDPXN20	нDQнM36
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		Thr-20 to Gly-26.
	628	659
	260 - 313	191 - 292
	124	125
	838139	801898
	HDTAU35	HDTAV54
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	1731 - 1754 6
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	838140	553651	753265
	не2сн58	HE2CM39	нЕ2НС60
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S0134: 1, H0483: 1, H0663: 1, L0005: 1, S0442: 1, S0354: 1, S0444: 1, S0442: 1, S0354: 1, S0444: 1, S0446: 1, L0008: 1, L0777: 1, H0411: 1, H0549: 1, L0024: 1, H0549: 1, H0634: 1, H0638: 1, H0634: 1, H0637: 1, H0647: 1, H0648: 1, H0634: 1, H0634: 1, H0648: 1, L0764: 1, L0549: 1, L0749:	L0756: 1, L0780: 1, L0757: 1, L0759: 1, S0242: 1 and S0456: 1.	AR089: 28, AR060: 14 H0556: 2, L0534: 2, L0562: 2, L0539: 2, L0109:
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		2, H0685: 1, S0114: 1	S0114: 1,	
		H0583: 1, HC	H0583: 1, H0657: 1, S0029:	
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		S0408: 1, H0	S0408: 1, H0619: 1, H0261:	-
		1, S0222: 1, H0587:	H0587: 1,	
		H0333: 1, S0	H0333: 1, S0280: 1, L0021:	
		1, H0098: 1, S0010:	S0010: 1,	
		H0052: 1, H0	H0052: 1, H0150: 1, H0172:	
		1, H0024: 1, T0010:	T0010: 1,	
		H0266: 1, S0	H0266: 1, S0003: 1, H0428:	
		1, H0070: 1, L0483:	L0483: 1,	
		H0030: 1, H0	H0030: 1, H0032: 1, H0316:	
		1, S0036; 1, H0090;	H0090: 1.	
	-	H0591: 1, H0	H0591: 1, H0372: 1, H0714:	
		1, H0646; 1, H0652;	H0652: 1.	
		L0598: 1, L0	L0598: 1, L0520: 1, L0762:	
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		L4747: 1, L5:	L4747: 1, L5565: 1, L0667:	
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		L0645: 1, L0	L0645: 1, L0764: 1, L0771:	
-		1, L0767: 1, L0533:	L0533: 1,	
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		1, L0606: 1, L0558:	L0558: 1,	
		L0809: 1, L0	L0809: 1, L0519: 1, L0647:	
		1, L0789: 1, L0664: 1,	L0664: 1,	
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		1, H0682: 1, H0659:	H0659: 1,	
		H0670: 1, L0	H0670: 1, L0602: 1, H0710:	
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		1, L0756: 1, L0780:	L0780: 1,	
		L0755: 1, S0 <sup>2</sup>	L0755: 1, S0434: 1, L0603:	
		1, S0011: 1, S0026: 1	S0026: 1,	
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	HE9CY05	HE9EA10
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		Val-40 to Cys-45, Lys-58 to Thr-64.	Arg-18 to Lys-26, Gly-35 to Ala-42, Gln-61 to Gly-67.
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	633719	831464	600355
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	HEBDF77	невр091	HEBFR46
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	960862	834379	693175	637624	674456	851137	834491	866171
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	169	170
	844543	790557
	HETCI16	HETDW58
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H05	5, L(	COTI	1,505	4, H	804	3, H	H04	3, H	HOI	3, L(	S10S	2. SC	804	2. SC	803	2.80	HO5	2. H	H00	2, H	00H H00	2, S(	T02	2, L	)T00	2, U	90H H06	2, H		2.8	2008	2, H	H03		H03
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L0005: 1. S0045: 1. H0619:	l, H0411: 1, H0175: 1,	H0369: 1, H0431: 1, H0392:	l, H0455: 1, H0612: 1,	H0587: 1, H0331: 1, L0622:	, H0486: 1, H0635: 1,	H0599; 1, H0098; 1, S0010;	, H0318: 1, H0310: 1,	H0263: 1, T0110: 1, H0545:	, N0006: 1, H0123: 1,	H0050: 1, H0011: 1, H0620:	T.0163: 1. T0010: 1.	H0083: 1, H0375: 1, S6028:	H0028 1 S0250 1	\$0214.1 H0328.1 H0039.	H0031-1 H0553-1	H0124: 1 H0598: 1 S0036:	H0038: 1 H0063: 1	TOOK7: 1 HOOK4: 1 HOA13:	1000/: 1, F0204: 1, F0413:	1, H0023: 1, 30038: 1, H0100: 1 1 0564: 1 H0043:	JO: 1, LO304: 1, 10042:	i, H0494: 1, H0625: 1,	H0561: 1, S0150: 1, L0598:	I, L0763: 1, L0761: 1,	.0667: 1, L0641: 1, L0650:	, L0375: 1, L0523: 1,	.0805: 1, L0654: 1, L0776:	, L0807: 1, L0647: 1,	.0792: 1, L0793: 1, L0666:	, L0664: 1, L0665: 1,	H0699: 1, S0374: 1, L0438:	, H0689: 1, H0435: 1,	H0659: 1, H0670: 1, H0660:	, L0602: 1, H0627: 1,	S0037: 1, S0027: 1, L0743:	l, L0749: 1, L0779: 1,	H0595: 1, L0605: 1, L0485:	, L0604: 1, L0593: 1,
0007	1, H0	H036	1, H0	H058	1, H0	H059	1, H0	H026	1, NO	H005	01.1	H008	1 H0	1208	1 HO	H012	TOTAL	T., T.	011 1	OH, II	HOIOT	1, H0	H056	1, L0	9907	1, L0	L080	1, L0	L079	1, 10	690H	1, H0	H065	1,10	500S	1, L0	H059	1, L0
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L0594: 1, S0196: 1 and S0412: 1.	AR089: 13, AR060: 9 H0305: 3, L0743: 3, H0620: 2, H0617: 2, L0770: 2, L0794: 2, L0384: 2, L0666: 2, L0777: 2, L0591: 2, L0595: 2, H0556: 1, S0358: 1, S0045: 1, H0497: 1, H0493: 1, H0618: 1, H0318: 1, H0581: 1, H0612: 1, H0428: 1, H0687: 1, L0351: 1, H0132: 1, H0529: 1, L0761: 1, L0644: 1, L0375: 1, L0524: 1, L0653: 1, L0659: 1, L0656: 1, L0659: 1, L0656: 1, H0650: 1, H0682: 1, H0670: 1, H0672: 1, H0555: 1, L0749: 1, L0779: 1, L0780: 1, L0749: 1, L0749: 1, H0653: 1, R0445: 1, H0653: 1, R0445: 1, H0653: 1, R0445: 1, H0653: 1, R0445: 1,	AR311: 142, AR272: 136, AR308: 126, AR104: 116, AR264: 98, AR212: 96, AR061: 75, AR055: 74, AR060: 74, AR201: 60, AR312: 34, AR197: 33, AR312: 34, AR197: 33, AR309: 25, AR096: 26, AR309: 25, AR053: 25, AR252: 24, AR213: 20, AR205: 15, AR245: 14,
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Met-1 to Leu-11, Val-13 to Lys-19, Thr-30 to Asp-39, Thr-49 to Gly-68, Ala-78 to Gly-111, Pro-140 to Thr-163, Ser-169 to Ser-185, Glu-197 to Lys-204, Lys-210 to Asp-215, Glu-220 to Ser-231, Ser-255 to Leu-266, Thr-269 to Asp-288, Cys-300 to Val-309, Phe-331 to Cys-339, Ser-362 to Ile-373.	
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	722235	799541
	HNEAK81	HNECL22
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*	800	801	802
	316 - 489	70 - 171	676 - 774
	296	297	298
	639117	815675	815676
	HNECW49	HNEDH88	HNFAC50
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	803	804
	314 - 445	178 - 270
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	825417	722237
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H0650: 1, H0657: 1, H0638: 1, S0354: 1, S0360: 1, S03408: 1, S0408: 1, S0408: 1, S0408: 1, S0408: 1, S0404: 1, H0441: 1, H0497: 1, H0331: 1, T0109: 1, S0278: 1, H0705: 1, H0318: 1, S0474: 1, T0110: 1, H0565: 1, H0572: 1, H0560: 1, H0573: 1, H0687: 1, S0408: 1, H0687: 1, S0448: 1, H0573: 1, L0741: 1, H0494: 1, L0772: 1, L0372: 1, L0800: 1, L0772: 1, L0372: 1, L0659: 1, L0774: 1, L0659: 1, L0774: 1, L0659: 1, L0664: 1, L0657: 1, L0659: 1, L0664: 1, S0428: 1, S0053: 1, H0444: 1, S0448: 1, H0648: 1, H0672: 1, L0439: 1, L0777: 1, L0439: 1, L0777: 1, L0749: 1, L0777: 1, L0763: 1, H0648: 1, H0627: 1, H0631: 1, S0027: 1, L0749: 1, L0777: 1, L0763: 1, L0749: 1, L0777: 1, L0763: 1, H0445: 1, S0462: 1, S0446: 1, S0462: 1, S0462	AR089: 26, AR060: 14 S0052: 1	AR089: 35, AR060: 16 S0052: 1
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	248 - 346	68 - 412
	301	302
	603910	688114
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	7, AR089: I		5, AR060:	4, AK254:	3, AR271:	3, AR061:	3, AR053:	3, AR201:	AR096:	AR249:	2, AK310: 2, AR265:	2, AR253:	, AR213:	l, AR273:	I, AR039:	1, AR205:	.0766: 3	2, H0402: 2, H0620:	0754: 2.	1, H0484: 1, H0254: 1,	0208: 1,	1, S0222: 1, H0618:	[0457: 1	1, H0051: 1, H0271: 1,	10063: 1	1, L0351: 1, T0042: 1,	S0448: 1, L0761: 1, L0378:	1, L0805: 1, L0655: 1,	H0539: 1, S0188: 1, S0146 1 H0543: 1 and H0423: 1.		
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	AR060: S0052	)OS	AR251:	AR052:	AD277	AR244:	AR198:	AR089:	AR264:	AR309:	AK311:	AR 197	AR104:	AR308:	AR194:	AR252:	SO —	2, H	00H	1, H	S03	1, S.	H01	1, H	HOH	1, [	S04	1, L	H05	<u>;</u> T	
13.				,00																											130.
Leu-93 o Gly-1	Glu-24	Arg-33	Ser-39,	Gin-61	.0 <b>3c</b> 1-1.																									Ser-39	to Gly-
Lys-83 to Leu-93, Pro-103 to Gly-113.	Asn-14 to Glu-24.	Pro-28 to Arg-33	Ser-21 to Ser-39,	Gln-45 to Gln-61	Cys-124 to 361-139.																									Ser-21 to Ser-39	Gin-45 to Gin-61, Cys-124 to Gly-130
	208	808	608						•								-			_					-					1017	
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	47 - 187	05 - 384	237 - 965																											231 - 629	
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	532614	825389	1145071																											866177	
	3H53	3038	X18	•		,																	_								
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	846148
	HSLHG78
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	Arg-28 to Arg-35.		Ser-6 to Ser-14.	Ala-17 to Thr-26, Gly-49 to Gln-62.	Pro-2 to Arg-7,
	903	904	905	906	200
	485 - 610	941 - 955	164 - 208	1508 - 1696	206 - 334
	399	400	401	402	403
	777861	784054	527221	635131	827058
7 1 2 2 1 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	HSLHXIS	HSNAP85	HSNAZ09	HSNBM34	HSOAH16
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	806	606																															
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	560726	853393																															
	HSQBF66	НЅОДО85															-																
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3, 1.0	H044	3, H0	H058	2, 80	S013	2, H0	H048	2. SQ	HO08	2. H0	H062	2. SO	H029	2, H0	H067	2. H0	80210	2, LO	1,080	2, S0.	H051	2, S0	T090	2, LO	L059	2, L0	H017	1, HO	S602	1, S02	1000S	1, S0	S044	1, H0	80278	1, HO	H036
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			406
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		2, H0545: 2, H0012: 2,	
		H0617: 2, H0135: 2, H0494:	
		2, L0763: 2, L0769: 2,	
 		L0768: 2, L0657: 2, L0438:	
		2, H0520: 2, H0658: 2,	
		H0539: 2, S0152: 2, L0747:	
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		L0588: 2, L0605: 2, S0040:	
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		S0360: 1. H0675: 1. H0645:	
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		H0550: 1, S6016: 1, H0370:	
		1 H0497 1 H0574 1	
		H0575-1 H0253-1 S0049-	
		1 T0115 1 H0530 1	
•		H0041: 1, H0620: 1, H0373:	
		1, H0375; 1, H0188; 1.	
		H0286: 1, H0181: 1, H0124:	
		1, H0068: 1, H0040: 1,	
 		H0412: 1, T0042: 1, H0561:	
-		1, S0448: 1, H0633: 1,	
		S0144: 1, S0210: 1, S0002:	
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		L0387: 1, L0375: 1, L0651:	
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		L0731: 1, L0757: 1, L0592:	

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	658725
	HSSEF77
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751308	895392	740766
HSWBE76	HSXCP38	HSYB106
403	404	405

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			Ala-15 to Gln-22, Gly-36 to Gly-41, Arg-47 to Pro-63, Pro-85 to His-98.
·	920	921	922
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	416	417	
	630647	838620	853400
	HT1SC27	HT3BF49	HT4FV41
	406	407	408

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590.1	3, HG	H025	2, S0	)00H	2, H(	990T	2, LC	7SOH	2. So	H043	JH I	1, 11	102	1, 50	8004	1, H(	H05	1. H(	750H	1. TC	H017	1, LC	H012	1, H(	H04	1, H(	H052	1, 10	L077	1, 10	790T	1, LC	T018	1, SC	90H	1, SC	L078	
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		975	976	776	978
	439 - 630	198 - 362	134 - 310	221 - 397	52 - 153
	470	471	472	473	474
	732808	853621	806212	762851	840596
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	460	461	462	463	464

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	Leu-12 to Cys-18.	Ser-83 to Asp-88, Val-166 to Gly-181, Pro-193 to Ala-199, Glu-235 to Gln-250.
	979	086
	93 - 149	84 - 884
	475	476
	637720	853401
	HTSFJ32	HTTCB60
	465	466

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S0045: 1, H0607: 1, H0586: 1, H0587: 1, T0040: 1, S0280: 1, H0590: 1, S0010: 1, H0581: 1, H0581: 1, H0581: 1, H0581: 1, H0641: 1, H0581: 1, H0651: 1, H0650: 1, H0681: 1, H06063: 1, H0612: 1, H0613: 1, L0761: 1, L0761	1 and H0352: 1, A0194: 1, H00008: 1 and H0352: 1.  AR089: 25, AR060: 17  H0052: 24, H0040: 17, L0768: 15, H0251: 14, L0769: 9, L0439: 9, L0770: 8, L0748: 8, L0731: 8, H0543: 8, H0423: 8, H0264: 7, H0494: 7, L076: 7, H0494: 7, L076: 7, L0659: 7, L0666: 7, H0144: 7, H0659: 7, L0669: 7, L0776: 7, L0777: 7, L0592: 7, S0222: 6, H0038: 6, H0529: 6, L0662: 6, H0435: 6, H0013: 5, H0318: 5, H0581: 5, H0012:
	H, 6, H, 7, C, 7, H, 8, C, C, F, A, F, B, F, F, C, C, F, E, C, C, F, E, C, C, E, C, E, C, E,
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5, H0616: 5, S0440: 5.	L0775: 5, L0663: 5, H0519:	8: 5, H0	4. S004	0: 4, H0	4, H054	9: 4, LO	4. H017	6.3 HC	2 5036	J, UV. V.	): 5, LU/	3, T011'	I: 3, HO	3 H008	2, 11000 7 7 TTO	5: 3, HU	3, H002'	3.3.50	7. 5, 5,	L0761: 3, L0375: 3, L0664:	9: 3, H0	3 H052	2, 11022	1: 3, 1:0	3, S043(	7: 2. H0	2, S041	5: 2. HO	7 H030	7, 1100/ 7. 0 UC	7. 2, 110	, HU25	3: 2, SOC	2, S000?	): 2, LO	2. H003	2.2.10	5003	1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1	7. 2, 19	., H064¢	: 2, LO¢	71.0377
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	Arg-23 to Leu-28.
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		4, L0749: 4, L0605: 4,		
		H0427: 3, H0050: 3, H0673:		
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		2, H0057: 2, H0014: 2,		
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		L0794: 2. L0766: 2. L0649:		
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		S0374-2 H0682-2 H0658-		
		20274: 2, 110082: 2, 110036: 2		
		2, HU321: 2, LU/42: 2,		
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		HU341: 1, HU459: 1, SU360:		
		1, S0046: 1, S0132: 1,		
		H0619: 1, H0392: 1, H0586:		
		1, T0109: 1, H0013: 1,		
		L0021: 1, H0599: 1, H0098:		
		1, H0575: 1, S0010: 1,		
		H0318: 1, S0474: 1, H0581:		
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		H0039: 1. H0553: 1. H0606:		
		1, S0366: 1, H0090: 1.		
		H0591: 1. H0616: 1. H0551:		
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	983	984
	66 - 311	350 - 388
	479	480
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4 HO	21: 4, I	: 4, LO	28: 4, L	: 4, LO7	42: 4, F	: 3, HO	18: 3, S	3, H05	86: 3, I	3, H03	31: 3, F	3, HO	)2: 3. S	3. L06	27: 3.1	3. H0	)6: 3. I	3.504	39:3	3. S01	22: 3, F	2, H02	)2: 2, F	2, S04	28: 2, S	2, H04	97: 2, H	2, H05	73: 2, S	2, H06	14: 2, H	2, H00	0: 2, S	2, L05	8: 2, L	2, L08
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L0475: 1, H0560: 1, H0625: 1, H0561: 1, S0438: 1, H0509: 1, H0132: 1, H0633: 1, H0650: 1, S0438: 1, H0650: 1, S0438: 1, H0650: 1, UNKWN: 1, L0520: 1, UNKWN: 1, L0550: 1, L0505: 1, L0505: 1, L0573: 1, L0372: 1, L0372: 1, L0373: 1, L0372: 1, L0373: 1, L0572: 1, L0378: 1, L0661: 1, L0807: 1, L0661: 1, L0807: 1, L0664: 1, S0428: 1, S0053: 1, H0713: 1, H0690: 1, L0590: 1, S0990: 1, H0590: 1, H0690: 1, H0590: 1, H0690: 1, H0590: 1, H0690:	AR253: 13, AR243: 12, AR254: 11, AR060: 10, AR250: 10, AR204: 9, AR061: 9, AR246: 9, AR205: 8, AR309: 8, AR039: 7, AR201: 7, AR089: 6, AR198: 6,
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1	AR272: 5, AR312: 5, AR272: 5, AR096: 5,		AR212: 4, AR308: 4,			AR213: 3, AR033: 2	L0755: 6, L0769: 4,	H0009: 3, H0012: 3, L0783:	3. L0749: 3. L0750: 3.	.9/£08 £ 18/01 £ .6/20 I	2. H0545: 2. S0051: 2.	H0606: 2, H0100: 2, 1,0638:	2. L0665; 2. S0028; 2.	L0751: 2. L0747: 2. L0756:	2. L0758; 2. L0603; 2.	H0265: 1, H0294: 1, H0341:	1, S0212: 1, S0410: 1,	S0045: 1, H0393: 1, H0549:	1, S0222: 1, H0586: 1,	H0331: 1, L0623: 1, T0060:	1, H0581: 1, S0049: 1,	H0309: 1, L0471: 1, H0620:	1, H0024: 1, H0266: 1,	H0428: 1, H0213: 1, L0456:	1, S0366: 1, S0036: 1,	H0040: 1, S0142: 1, L0763:	1, L0371: 1, L0772: 1,	L0372: 1, L0646: 1, L0764:	1, L0773: 1, L0766: 1,	L0774: 1, L0775: 1, L0776:	1, L0809: 1, L0519: 1,	H0520: 1, H0519: 1, S0126:	1, H0660: 1, S0350: 1,	S0406: 1, H0436: 1, S3012	1, L0752: 1, L0757: 1,	S0436: 1, L0592: 1, S0276
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1 and H0422: 1.	AR089: 168, AR060: 110 1.0659: 33, 1.0665: 27	L0666: 19, L0664: 19,	S0360: 17, S0344: 17, 1.0648: 17, S0358: 16	L0655: 14, L0596: 13,	L0751: 12, L0662: 11,	L0663: 10, L0740: 9, L0775:	8, L0599: 8, S0376: 7,	H0046: 7, H0486: 6, H0597:	6, S0126: 6, L0439: 6,	L0752: 6, S0116: 5, S0140:	5, H0581: 5, S0328: 5,	L0748: 5, H0543: 5, H0423:	5, H0657: 4, S0212: 4,	H0617: 4, H0087: 4, S0372:	4, L0374: 4, L0651: 4,	H0555: 4, L0744: 4, L0754:	4, L0747: 4, T0049: 3,	S0278: 3, H0031: 3, H0641:	3, S0144: 3, L0646: 3,	L0375: 3, L0776: 3, L0606:	3, L0661: 3, L0657: 3,	S0428: 3, H0518: 3, H0521:	3, S3014: 3, L0742: 3,	L0743: 3, L0750: 3, L0753:	3, L0362: 3, L0601: 3,	S0026: 3, H0265: 2, H0556:	2, T0002: 2, H0686: 2,	S0114: 2, H0402: 2, S0410:	2, S0300: 2, T0060: 2,	H0575: 2, H0274: 2, H0318:	2, H0085: 2, H0231: 2,	H0083: 2, H0271: 2, H0188:	2, H0688: 2, H0553: 2,	H0068: 2, H0509: 2, S0142:	2, S0002: 2, L0369: 2,
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	Prò-70 to Ser-89, Ser-92 to Ser-115.	
	888	686
	49 - 447	216 - 416
	484	485
	853408	658730
	HTXDD61	HTXDG92
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54.8	7. C0	47: 5,	. 4, LO	65: 3,	: 3, HOC	17: 3,	3, L08	49: 3,	2. HO	16: 2.	., HOC	50.0	 	: 2, 1W	71: 2,	2. L07	00.5	7. HO5	50.0	39: 4,	2, H05	17: 1,	1, T00	56: 1,	: 1, S02	07: 1,	1, S02	50: 1,	: 1, H02	0: 1,	: 1, HO	57: 1,	: 1, H0C	94: 1,	: 1, H07	53: 1,	1, H0	63: 1,
1 0777 11 1 0754 8	1, 207.	6, H0144: 5, L0747: 5,	H0556: 4, H0059: 4, L0439:	4, L0601: 4, H0265: 3	S0410: 3, H0253: 3, H0052:	3, H0620: 3, H0617: 3	L0764: 3, L0768: 3, L0806:	3, L0744: 3, L0749: 3,	L0731: 3, H0583: 2, H0341:	2, S0358; 2, S0046; 2,	S0222: 2, H0013: 2, H0069-	2. S0049: 2, H0150: 2,	1.0251	HUU8/: 2, LU351: 2, 10041:	2, H0529: 2, L0771: 2	L0773: 2, L0662: 2, L0766:	2, L0803; 2, L0809; 2	1,0793 2 1,0665 2 H0547	20173: 2, 20003: 2, III 2 H0510: 2 H0650: 2	2, EG.	L0742: 2, L0748: 2, H0542:	2, S0040: 1, H0717: ]	H0716: 1, S0114: 1, T0049:	1, H0657: 1, H0656:	H0381: 1, H0663: 1, S0360:	1, H0722: 1, S0007: 1	S0132: 1, S0300: 1, S0278:	1, H0261: 1, H0550: 1	H0431: 1, H0392: 1, H0486:	1, T0114: 1, S0010: 1	H0581: 1, H0374: 1, H0327:	, H0545: 1, H0457: 1	H0012: 1, H0024: 1, H0015	, H0510: 1, H0594: 1,	H0188: 1, H0292: 1, H0286:	, H0622: 1, H0553: 1	H0181: 1, H0135: 1, H0163:	1, H0040: 1, H0063: 1, H0413: 1 H0100: 1 H0561:
.7770	0618:7	H0144	3556: 4	L0601:	410:3,	H0620:	1764: 3,	L0744:	1731: 3.	\$0358	0000	S0049.		1007: 2,	H0529:	773: 2,	1.0803:	793.2	U0510	71000	742: 2,	S0040:	716: 1,	H0657:	J381: 1,	H0722:	132: 1,	H0261:	431:1,	Γ0114:	581: 1,	H0545:	012: 1,	H0510:	188: 1,	H0622:	181: 1,	H0040:
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				Met-1 to Pro-6, Gly-73 to Thr-78.
	066	991	992	993
	178 - 267	192 - 281	108 - 158	330 - 566
	486	487	488	489
	581521	853410	637774	834438
	HTXET11	HTXFA72	HTXJY08	HTXKF95
	476	477	478	479

L0754: 41, L0747: 8, L0755: 5, L0659: 4, H0265: 2, H0556: 2, H0586: 2, L0471: 2, H0553: 2, L0764: 2, L0662: 2, L0794: 2, L0748: 2, L0751: 2, L0749: 2, L0750: 2, H0305: 1, S0358: 1, S0046: 1, H0441: 1, H0599: 1, H0569: 1, H0050: 1, H0616: 1, L0770: 1, L0769: 1, L0800: 1, L0644: 1, L0363: 1, L0806: 1, L0783: 1, L0806: 1, L0665: 1, H0144: 1, L0806: 1, L0773: 1, L0779: 1, L0773: 1, L0779: 1, L0773: 1, L0666: 1, L0653: 1, L0666: 1, L0665: 1, L0779: 1, L0731: 1, L0665: 1, L0779: 1, L0731: 1, L0779: 1, L0779: 1, L0779: 1, L0731: 1, L0779: 1, L0779: 1, L0731: 1, L0779: 1, L0779	AR060: 7, AR089: 4 L0439: 6, H0556: 2, S0007: 2, L0744: 2, L0740: 2, L0731: 2, S0442: 1, L0021: 1, H0618: 1, H0253: 1, H0041: 1, L0770: 1, L0800: 1, L0766: 1, L0803: 1, L0375: 1, L0807: 1, L0382: 1, L0791: 1, L0793: 1, L0352: 1, S0432: 1, L0741: 1 and L0779: 1.	AR060: 26, AR089: 7 L0764: 5, L0771: 5, H0506: 4, L0374: 3, S0434: 3, S0356: 1, S0408: 1, H0264: 1, L0372: 1, L0783: 1, L0532: 1 and L0663: 1.
·	Pro-19 to Ser-28.	
	994	995
	319 - 432	287 - 367
	490	491
	834881	801938
	HTXMZ07	HUFCL31
	480	481

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AR089: 14, AR060: 9	H0052: 13, S0360: 8,	L0748: 8, H0619: 6, L0659:	6, L0665: 6, L0759: 6,	L0789: 5, L0743: 5, L0752:	5, S0346: 4, H0059: 4,	L0662: 4, L0805: 4, H0521:	4, L0717: 3, H0599: 3,	H0644: 3, L0761: 3, L0776:	3, S0028: 3, L0744: 3,	L0754: 3, L0749: 3, L0731:	3, L0757: 3, S0001: 2,	S0354: 2, H0261: 2, H0586:	2, S0010: 2, H0620: 2,	L0771: 2, L0804: 2, L0774:	2, L0806: 2, L0809: 2,	L0664: 2, H0547: 2, H0539:	2, H0555: 2, L0747: 2,	L0750: 2, L0758: 2, S0434:	2, L0596: 2, L0604: 2,	H0171: 1, S0040: 1, H0713:	1, H0656: 1, S0212: 1,	L0005: 1, S0356: 1, H0728:	1, H0733: 1, S0046: 1,	S0278: 1, H0370: 1, H0392:	1, H0602: 1, H0592: 1,	H0574: 1, H0013: 1, S0280:	1, H0575: 1, T0082: 1,	H0581: 1, H0544: 1, H0046:	1, H0009: 1, H0081: 1,	H0051: 1, H0266: 1, H0179:	1, H0290: 1, H0286: 1,	S0250: 1, S0366: 1, S0036:	1, H0135: 1, H0591: 1,	H0038: 1, H0551: 1, H0264:	1, H0488: 1, T0004: 1,	H0100: 1, H0429: 1, H0334:	1, H0386: 1, S0144: 1,
Ser-32 to Arg-39.	•											,																									
966																																					
273 - 392																										-											
492																																					
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HUKBT67											•								_																		
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## DOGGOON TOOLOGE

S0344: 1, S0002: 1, L0763: 1, L0667: 1, L0764: 1, L0773: 1, L0794: 1, L0766: 1, L0803: 1, L0650: 1, L0657: 1, L0793: 1, L0666: 1, S0053: 1, H0144: 1, L0352: 1, H0520: 1, H0660: 1, H0672: 1, S0328: 1, H0696: 1, S0404: 1, S0406: 1, H0436: 1, S0328: 1, S0037: 1, L0742: 1, L0779: 1, L0777: 1, S0031: 1, S0260: 1, L0584: 1, L0591: 1 and H0506: 1.	IR060: 6, AR089: 3 H0266: 1 and H0059: 1.	AR089: 14, AR060: 8 S0053: 4, H0673: 3, H0618: 2, H0179: 2, H0674: 2, S0216: 2, H0521: 2, S0031: 2, H0556: 1, S0116: 1, H0305: 1, H0619: 1, H0550: 1, H0609: 1, H0635: 1, H0318: 1, H0309: 1, H0083: 1, H0271: 1, H0090: 1, H0634: 1, H0059: 1, S0002: 1, S0428: 1, H0144: 1, S0152: 1 and L0740: 1.	5, AR194: 4, 3, AR251: 3, 2, AR205: 2, 2, AR039: 2, 2, AR060: 1, 1, AR311: 1, 1, AR186: 1, 1, AR263: 1
S0344: 1, S0002: 1, I 1, L0667: 1, L0764: 1 L0773: 1, L0794: 1, I 1, L0803: 1, L0500: 1 L0657: 1, L0793: 1, I 1, S0053: 1, H0144: 1 L0352: 1, H0520: 1, I 1, H0672: 1, S0328: 1 H0696: 1, S0404: 1, S 1, H0436: 1, S0404: 1, S 1, H0436: 1, S0402: 1, L0777: 1, S031: 1 S0260: 1, L0742: 1, I 1 L0777: 1, S0031: 1 S0260: 1, L0584: 1, I I and H0506: 1.	AR060: 6, AR089: H0266: 1 and H005	AR089: 14, AR060: 8 S0053: 4, H0673: 3, H0618: 2, H0179: 2, H0 2, S0216: 2, H0521: 2, S0031: 2, H0556: 1, S0 1, H0305: 1, H0619: 1, H0550: 1, H0609: 1, H0 1, H0318: 1, H0309: 1, H0083: 1, H0271: 1, H0 1, H0634: 1, H0059: 1, S0002: 1, S0052: 1, S04 1, H0144: 1, S0152: 1 a L0740: 1.	AR245: 5, AR061: 3, AR201: 2, AR198: 2, AR055: 2, AR204: 2, AR312: 1, AR243: 1,
			Phe-166 to Arg-174, Ser-191 to Tyr-196.
	266	866	666
	214 - 315	187 - 285	74 - 661
	493	494	495
	566823	570896	894699
	HUKDF20	HUKDY82	HUSCI14
	483	48	88

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L0777: 8, L0766: 7, L0741:	L0754: 5, L0744: 4, L0757:	4, S0192: 4, H0677: 4,	H0556: 3, S0360: 3, S0410:	3, H0013: 3, H0052: 3,	L0769: 3, L0775: 3, L0776:	3, L0756: 3, L0752: 3,	L0604: 3, H0265: 2, S0040:	2, H0599: 2, H0545: 2,	H0266: 2, H0030: 2, H0617:	2, H0135: 2, L0771: 2,	L0662: 2, L0806: 2, L0805:	2, L0659: 2, L0666: 2,	L0665: 2, H0520: 2, H0547:	2, H0519: 2, H0659: 2,	S0404: 2, L0743: 2, L0758:	2, L0596: 2, L0605: 2,	L0485: 2, H0171: 1, H0713:	1, S0134: 1, S0218: 1,	H0657: 1, H0656: 1, S0212:	1, H0663: 1, S0420: 1,	S0408: 1, S0132: 1, S0476:	1, H0393: 1, H0587: 1,	T0040: 1, T0060: 1, H0575:	1, H0309: 1, H0009: 1,	L0471: 1, H0620: 1, H0510:	1, H0290: 1, S0250: 1,	S0022: 1, T0023: 1, L0055:	1, H0634: 1, H0488: 1,	H0268: 1, T0041: 1, T0042:	1, H0538: 1, S0210: 1,	L0763: 1, L0639: 1, L0764:	1, L0794: 1, L0649: 1,	L0804: 1, L0650: 1, L0774:	1, L0809: 1, L0793: 1,	L0664: 1, H0144: 1, H0593:
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H0521: 1, S0406: 1, H0555: 1, L0740: 1, L0747: 1, L0749: 1, L0779: 1, L0731: 1, L0759: 1, S0031: 1, S0434: 1, S0436: 1, H0665: 1, H0667: 1 and S0276: 1.	AR252: 82, AR250: 77, AR253: 70, AR254: 37, AR309: 22, AR264: 17, AR308: 16, AR312: 15, AR263: 15, AR096: 13, AR211: 11, AR271: 10, AR213: 8, AR243: 7, AR245: 7, AR053: 7, AR246: 6, AR272: 6, AR069: 6, AR272: 6, AR069: 6, AR212: 6, AR069: 6, AR212: 6, AR069: 3, AR033: 4, AR204: 4, AR033: 4, AR204: 4, AR033: 4, AR204: 4, AR033: 2, AR061: 3, AR104: 3, AR205: 2, L0766: 4, S0358: 3, H0266: 3, S0356: 2, H0616: 2, L0794: 2, L0655: 2, H0672: 2, L0777: 2, L0731: 2, H0483: 1, H0499: 1, S0360: 1, H0587: 1, H0497: 1, H0486: 1, H050: 1, R0377: 1, H0486: 1, H021: 1, H0377: 1, S6028: 1, H0375: 1, S6028: 1, H0328: 1, T0006: 1, H0644: 1, H0328: 1,
	Met-1 to Tyr-8,  Glin-27 to Glin-38.
	0001
	350 - 493
	496
	792637
	HUSGL67
	486

## DOGECONE, COTECT

1, H0090: 1, H0038: 1, H0087: 1, H0264: 1, H0268: 1, H0412: 1, S0422: 1, H0529: 1, L0521: 1, L0803: 1, L0659: 1, L0666: 1, H0710: 1, H0518: 1, H0521: 1, S0176: 1, S0406: 1, S3014: 1, L0439: 1, L0758: 1 and H0543: 1.	AR089: 34, AR060: 23	L0748: 4, H0622: 3, L0777: 3, H0624: 2, H0013: 2, H0520: 2, H0539: 2, L0439: 2, L0754: 2, L0747: 2, L0757: 2, L0758: 2, L0593: 2, L0002: 1, H0664: 1, H0580: 1, S0007: 1, H0497: 1, H0333: 1, H0599: 1, H0581: 1, L0483: 1, H0598: 1, L0748: 1, L0769: 1, L0771: 1, L0662: 1, L0767: 1, L0768: 1, L0766: 1, L0768: 1, L0766: 1, L0768: 1, L0766: 1, L078: 1, L0769: 1, L0769: 1, L0740: 1, L0759: 1, L0740: 1, L0752: 1, L0740: 1, L0752: 1, L0740: 1, L0752: 1,	AR060: 5, AR089: 3 H0393: 1, H0056: 1 and L0662: 1.	AR089: 4, AR060: 2 H0547: 12, L0794: 10, H0251: 9, L0439: 8, L0731:
	Arg-21 to Ser-27, Ile-36 to Asp-41.			
	1001	1002	1003	1004
	500 - 640	83 - 151	196 - 213	223 - 312
	497	498	499	500
	684975	762858	564853	830432
	HUSGU40	HUSIR18	HUVDJ48	HWAAI12
	487	488	489	490

		\$ 10747.7 10439.6	
	_	o, LU/4/. /, LU45o. U,	
		H0351: 5, L0750: 5, S0356:	
		4, L0768: 4, L0766: 4,	
		L0805: 4, L0809: 4, L0777:	
		4, L0758: 4, L0596: 4,	
	•	S0410: 3, H0013: 3, H0009:	
		3, H0594: 3, T0006: 3,	
		H0124: 3, T0041: 3, H0529:	
		3, L0769: 3, L0666: 3,	
		H0144: 3, H0520: 3, L0749:	
		3. H0661: 2. H0305: 2.	
		S0360: 2 10103: 2 H0591:	
		30300: 2, L0103: 2, H0301:	
		2, S0049: 2, H0052: 2,	
		L0157: 2, L0471: 2, H0031:	
		2, H0087: 2, H0100: 2,	
	• 44	S0344: 2, L0763: 2, L0761:	
		2, L0662; 2, L0803; 2.	
		L0806: 2, L0664: 2, H0436:	
		2. S0028: 2. 1.0742: 2.	
		L0756: 2, L0752: 2, L0605:	
		2, L0595; 2, H0543; 2.	
		H0556: 1, T0002: 1, S0040:	
		1, H0717: 1, H0716: 1,	
		H0294: 1, S0134: 1, H0341:	
-		1, H0402: 1, S0354: 1,	
		L0717: 1, S0278: 1, L0394:	
		1, H0549: 1, S0222: 1,	
		H0333: 1, L0622: 1, H0486:	
		1, H0250: 1, H0635: 1,	
		H0575: 1, S0010: 1, H0421:	
		1, H0085: 1, H0597: 1,	
		H0545: 1, H0566: 1, H0620:	
		1, H0271: 1, H0687: 1,	
		H0615: 1, H0622: 1, H0673:	
		1, H0674: 1, H0412: 1,	
		H0413: 1, H0056: 1, T0042:	
		1, H0130: 1, H0646: 1,	
		S0144: 1, S0208: 1, L0770:	

			,
1, L0796: 1, L0667: 1, L0772: 1, L0373: 1, L0372: 1, L0800: 1, L0645: 1, L0764: 1, L0648: 1, L0767: 1, L0650: 1, L0657: 1, L0517: 1, L0789: 1, L0790: 1, L0665: 1, H0690: 1, H0658: 1, H0670: 1, H0672: 1, S0378: 1, S0380: 1, H0521: 1, S3012: 1, S0390: 1, S0027: 1, L0743: 1, L0779: 1, L0755: 1, L0759: 1, S0031: 1, S0436: 1, L0601: 1, S0276: 1, H0542: 1 and S0424: 1.	AR060: 2, AR089: 1 L0717: 2, H0580: 1, S0222: 1, L0662: 1, H0436: 1, L0748: 1, H0445: 1 and S0308: 1.	AR060: 1, AR089: 1 H0580: 1	AR089: 11, AR060: 5 H0635: 7, L0794: 6, H0556: 4, S0414: 4, H0521: 4, H0634: 3, L0779: 3, H0265: 2, S0134: 2, S0360: 2, H0619: 2, H0069: 2, H0575: 2, H0688: 2, H0056: 2, S0002: 2, L0665: 2, S0216: 2, H0519: 2, L0751: 2, L0758: 2, L0593: 2, H0422: 2, S0114: 1, S0116: 1, H0300: 1, S0356: 1, H0581: 1, S0045: 1, S0046: 1, H0643: 1, H0250: 1, H0581: 1, S0049: 1, L0045:
	Ala-21 to Ser-31.	Lys-45 to Pro-51, Arg-80 to Arg-85.	Ser-30 to Gly-36.
·	1005	1006	1007
	222 - 353	378 - 650	253 - 405
	501	502	503
	689121	722259	762860
	нwввQ70	HWBCN36	HWBDJ08
	491	492	493

H0644: 1, H0551: 1, H0264: 1, H0623: 1, H0641: 1, H0646: 1, L0763: 1, L0536: 1, L0765: 1, L0653: 1, L0655: 1, H0134: 1, L0777: 1, L0755: 1, H0542: 1, H0543: 1 and H0543: 1.	AR060: 184, AR089: 165 S0114: 1 and H0580: 1.	AR089: 61, AR060: 49, AR198: 5, AR194: 4, AR096: 3, AR310: 3, AR265: 3, AR213: 2, AR312: 2, AR249: 2, AR312: 2, AR049: 2, AR052: 2, AR104: 2, AR205: 1, AR039: 1 H0580: 1, S0300: 1, H0600: 1, L0783: 1, L0438: 1, L0439: 1 and L0758: 1.	AR060: 28, AR089: 14 H0556: 19, H0265: 15, S0418: 10, S0358: 9, S0440: 9, L0755: 9, S0420: 8, L0752: 7, H0253: 6, L0751: 6, L0747: 6, L0750: 6, L0596: 6, S0212: 5, H0618: 5, H0545: 5, H0012: 5, H0617: 5, H0413: 5, L0740: 5, L0601: 5, H0295: 4, S0360: 4, H0039: 4, H0494: 4, H0641: 4, L0764: 4, L0776: 4, S0406: 4, L0758: 4, H0445: 4, H0657: 3, H0483: 3, S0356: 3, S0376: 3, S0408: 3, S0344: 3, L0637: 3, H0547: 3, H0658: 3,
	1008	1009	1010
	267 - 278	242 - 349	866 - 964
	504	505	906
	827312	821335	796743
	HWBFX16	HWDAC26	HWDAG96
	494	495	496

		110660: 3 50378: 3 110573.	
		110000. 3, 30326. 3, 110322.	
		3, L0/43: 3, L0/49: 3,	
		L0756: 3, L0731: 3, L0757:	
		3, S0040: 2, H0713: 2,	
		H0294: 2, H0341: 2, H0484:	
		2, H0661: 2, H0305: 2,	
 ·		H0125: 2, H0580: 2, H0586:	
 		2, H0587: 2, H0052: 2,	
		H0046: 2, H0009: 2, H0081:	
		2. H0620: 2. H0266: 2.	
<del></del>		H0124: 2, H0135: 2, H0551:	
		2 H0100: 2 I 0646: 2	
		1.0768.2 1.074.2 1.0806.	
		2 H0435: 2 H0530: 2	
		1,110433. 2,110333. 2,	
		LU/46: 2, LU/34: 2, LU366:	
		2, L0589: 2, L0608: 2,	
		L0593: 2, H0543: 2, S0384:	
		2, H0170: 1, H0140: 1,	
		H0716: 1, H0650: 1, H0656:	
		1, H0254: 1, H0300: 1,	
		H0638: 1, S0410: 1, H0637:	
		1, S0045: 1, S0046: 1,	
		S0476: 1, H0619: 1, S0278:	
		1, S0222: 1, H0600: 1,	
•		H0497: 1, H0632: 1, H0559:	
		1, H0013: 1, H0069: 1,	
		H0042: 1, H0706: 1, S0010:	
		1, S0182: 1, H0318: 1,	
		H0263: 1, T0110: 1, L0471:	
		1, H0024: 1, H0416: 1,	
		H0290: 1, H0292: 1, H0286:	
		1, S0250: 1, H0622: 1,	
		L0194: 1, L0483: 1, T0006:	
		1, H0213: 1, H0644: 1,	
		L0142: 1, H0181: 1, H0606:	
		1, L0055: 1, H0090: 1,	
		H0038: 1, H0616: 1, T0067:	
		1, H0488: 1, H0412: 1,	

H0056: 1, T0041: 1, T0042: 1, L0475: 1, H0396: 1, S0144: 1, S0142: 1, S0210: 1, S0002: 1, H0695: 1, L0763: 1, L0763: 1, L0763: 1, L0662: 1, L0649: 1, L0381: 1, L0388: 1, L0573: 1, L0570: 1, L0570: 1, L0570: 1, L0570: 1, L0571: 1, L0570: 1, L0571: 1, L0570: 1, L0571: 1, L0570: 1, H0698: 1, H0698: 1, H0698: 1, H0698: 1, H0672: 1, S0044: 1, H0672: 1, S0044: 1, H0672: 1, S0044: 1, H0672: 1, S0044: 1, H0679: 1, S0028: 1, L0590: 1, L0759: 1, L0759: 1, L0590: 1, L0759: 1, S0026: 1, H0521: 1, S0026: 1, H0521: 1, S0026: 1, H0590: 1, S0026: 1, H0520:	AR060: 2 H0600: 1	H0437: 2, H0587: 2, H0494: 2, L0769: 2, H0547: 2, S0028: 2, L0439: 2, L0593: 2, H0556: 1, H0657: 1, H0662: 1, H0125: 1, S0418: 1, H0619: 1, H0618: 1, H0253: 1, H0318: 1, H0052: 1, H0135: 1, H0529: 1, L0438: 1, H0539:
	Pro-17 to Ser-24.	Gln-25 to Leu-30.
	1011	1012
	288 - 362	200 - 400
	207	508
	794016	740778
	HWDAJ01	HWHPB78
	497	498

1, H0521: 1, S0037: 1, S0424: 1, H0506: 1 and H0008: 1	AR089: 10, AR060: 6 L0655: 10, L0754: 6, L0438: 5, L0751: 5, L0777: 5, L0752: 5, L0755: 4, L0758: 4, S0046: 3, H0213: 3, L0769: 3, L0667: 3, L0771: 3, L0662: 3, L0659: 3, H0539: 3, L0747: 3, L0757: 3, S0276: 3, S0418: 2, H0428: 2, H0424: 2, H0553: 2, H0412: 2, L0649: 2, L0764: 2, L0768: 2, L0649: 2, L0766: 2, L0668: 2, L0764: 2, L0768: 2, L0768: 2, L0699: 2, S0040: 1, S0342: 1, T0049: 1, H0580: 1, H0550: 1, H0570: 1, H0580: 1, S0132: 1, H051: 1, H0550: 1, H0370: 1, H0550: 1, H0099: 1, H0408: 1, H0650: 1, H0179: 1, H0211: 1, H0500: 1, H0059: 1, H0641: 1, S0210: 1, H0529: 1, L0639: 1, L0637: 1, H0529: 1, L0639: 1, L0637: 1,
	Pro-3 to Ala-8.
	1013
	1015 - 1203
	206
	789854
	HYABC84
	466

1, L0761: 1, L0646: 1, L0643: 1, L0773: 1, L0650: 1, L0657: 1, L0635: 1, L0383: 1, L0790: 1, L0792: 1, L0664: 1, S0052: 1, H0691: 1, H0593: 1, H0435: 1, H0672: 1, H0696: 1, H0576: 1, L0748: 1, L0745: 1, L0750: 1, L0731: 1, H0707: 1, L0596: 1, L0591: 1, L0592: 1, L0593: 1, L0595: 1, H0667: 1, H0422: 1 and L0600: 1.	AR089: 10, AR060: 6 L0665: 10, L0754: 6, L0438: 5, L0751: 5, L0777: 5, L0752: 4, L0755: 4, L0758: 4, S0046: 3, H0213: 3, L0769: 3, L0667: 3, L0771: 3, L0662: 3, L0659: 3, H0539: 3, L0747: 3, L0757: 3, S0276: 3, S0418: 2, H0208: 2, S0045: 2, H0428: 2, H0424: 2, H0553: 2, L0666: 2, L0688: 2, L0764: 2, L0688: 2, L0649: 2, L0666: 2, L0668: 2, L0439: 2, L0756: 2, L0439: 2, L0756: 2, L0439: 2, L0756: 2, L0485: 1, H0580: 1, S0132: 1, H0583: 1, H0657: 1, H0370: 1, H0586: 1, H0333: 1, H0013: 1, H0550: 1, S0280: 1, H0575: 1, H0618:
	Pro-3 to Ala-8.
	8 1014
	1080 - 1268
	510
	865064
	HYABC84
	200

H																						
	H0009: 1, L0471: 1, H0620:		10408:	;	H0179: 1, H0271: 1, H0124:	٠.	10509:		H0529: 1, L0639: 1, L0637:		0650:	1, L0657: 1, L0635: 1,	0792:		10435:		.0745:		.0591:		L0595: 1, H0667: 1, H0422:	
052: 1	71: 1, F	388: 1	0: 1, H	0266: 1	71: 1, I	1135: 1	12: 1, F	)210: 1	39: 1, L	1946: 1	3: 1, L	635: 1	0: 1, L	052: 1	93: 1, F	3696: 1	1, I	731: 1	6: 1, I	593: 1	57: 1, E	
): 1, H(	1, L04	3: 1, SC	I, T001	9: 1, H	1, H02	5: 1, H(	1, T00 <sup>2</sup>	1: 1, SC	1, L06	I: 1, LC	1, L077	7: 1, LC	1, L079	t: 1, S0	1, H05	2: 1, H(	1, L074	): 1, L0	1, L'059	2: 1, L0	, H066	600: 1
1, S0049: 1, H0052: 1,	:0000	, L016.	0051:	, H023	10179:	, \$0366	[0059:	, H064	10529:	, L076	L0643: 1, L0773: 1, L0650:	,L0657	0383:	, L066	0691:	H067	.0576:	1, L0750: 1, L0731: 1,	H0707: 1, L0596: 1, L0591:	L0592	0595: 1	and L0600: 1.
_	<u></u>		S	1	正	_	<u> </u>		<u> </u>		<u>1</u>		7	1	王		<u> </u>	<u> </u>	正	<u> </u>	<u> </u>	_
																		-				
	_							_							,							
					_			_												_		

- [81] The first column in Table 1B provides the gene number in the application corresponding to the clone identifier. The second column in Table 1B provides a unique "Clone ID NO:Z" for a cDNA clone related to each contig sequence disclosed in Table 1B. This clone ID references the cDNA clone which contains at least the 5' most sequence of the assembled contig and at least a portion of SEQ ID NO:X was determined by directly sequencing the referenced clone. The reference clone may have more sequence than described in the sequence listing or the clone may have less. In the vast majority of cases, however, the clone is believed to encode a full-length polypeptide. In the case where a clone is not full-length, a full-length cDNA can be obtained by methods described elsewhere herein.
- [82] The third column in Table 1B provides a unique "Contig ID" identification for each contig sequence. The fourth column provides the "SEQ ID NO:" identifier for each of the contig polynucleotide sequences disclosed in Table 1B. The fifth column, "ORF (From-To)", provides the location (i.e., nucleotide position numbers) within the polynucleotide sequence "SEQ ID NO:X" that delineate the preferred open reading frame (ORF) shown in the sequence listing and referenced in Table 1B, column 6, as SEQ ID NO:Y. Where the nucleotide position number "To" is lower than the nucleotide position number "From", the preferred ORF is the reverse complement of the referenced polynucleotide sequence.
- [83] The sixth column in Table 1B provides the corresponding SEQ ID NO:Y for the polypeptide sequence encoded by the preferred ORF delineated in column 5. In one embodiment, the invention provides an amino acid sequence comprising, or alternatively consisting of, a polypeptide encoded by the portion of SEQ ID NO:X delineated by "ORF (From-To)". Also provided are polynucleotides encoding such amino acid sequences and the complementary strand thereto.
- [84] Column 7 in Table 1B lists residues comprising epitopes contained in the polypeptides encoded by the preferred ORF (SEQ ID NO:Y), as predicted using the algorithm of Jameson and Wolf, (1988) Comp. Appl. Biosci. 4:181-186. The Jameson-

Wolf antigenic analysis was performed using the computer program PROTEAN (Version 3.11 for the Power MacIntosh, DNASTAR, Inc., 1228 South Park Street Madison, WI). In specific embodiments, polypeptides of the invention comprise, or alternatively consist of, at least one, two, three, four, five or more of the predicted epitopes as described in Table 1B. It will be appreciated that depending on the analytical criteria used to predict antigenic determinants, the exact address of the determinant may vary slightly.

[85] Column 8, in Table 1B, provides an expression profile and library code: count for each of the contig sequences (SEQ ID NO:X) disclosed in Table 1B, which can routinely be combined with the information provided in Table 4 and used to determine the tissues, cells, and/or cell line libraries which predominantly express the polynucleotides of the invention. The first number in column 8 (preceding the colon), represents the tissue/cell source identifier code corresponding to the code and description provided in Table 4. For those identifier codes in which the first two letters are not "AR", the second number in column 8 (following the colon) represents the number of times a sequence corresponding to the reference polynucleotide sequence was identified in the tissue/cell source. Those tissue/cell source identifier codes in which the first two letters are "AR" designate information generated using DNA array technology. Utilizing this technology, cDNAs were amplified by PCR and then transferred, in duplicate, onto the array. Gene expression was assayed through hybridization of first strand cDNA probes to the DNA array, cDNA probes were generated from total RNA extracted from a variety of different tissues and cell lines. Probe synthesis was performed in the presence of <sup>33</sup>P dCTP, using oligo(dT) to prime reverse transcription. After hybridization, high stringency washing conditions were employed to remove non-specific hybrids from the array. The remaining signal, emanating from each gene target, was measured using a Phosphorimager. Gene expression was reported as Phosphor Stimulating Luminescence (PSL) which reflects the level of phosphor signal generated from the probe hybridized to each of the gene targets represented on the array. A local background signal subtraction was performed before the total signal generated from each array was used to normalize gene expression between the different hybridizations. The value presented after "[array code]:" represents the mean of the duplicate values, following background subtraction and probe normalization. One of skill in the art could routinely use this information to identify normal and/or diseased tissue(s) which show a predominant expression pattern of the corresponding

polynucleotide of the invention or to identify polynucleotides which show predominant and/or specific tissue and/or cell expression.

[86] Column 9 in Table 1B provides a chromosomal map location for certain polynucleotides of the invention. Chromosomal location was determined by finding exact matches to EST and cDNA sequences contained in the NCBI (National Center for Biotechnology Information) UniGene database. Each sequence in the UniGene database is assigned to a "cluster"; all of the ESTs, cDNAs, and STSs in a cluster are believed to be derived from a single gene. Chromosomal mapping data is often available for one or more sequence(s) in a UniGene cluster; this data (if consistent) is then applied to the cluster as a whole. Thus, it is possible to infer the chromosomal location of a new polynucleotide sequence by determining its identity with a mapped UniGene cluster.

[87] A modified version of the computer program BLASTN (Altshul, et al., J. Mol. Biol. 215:403-410 (1990), and Gish, and States, Nat. Genet. 3:266-272) (1993) was used to search the UniGene database for EST or cDNA sequences that contain exact or near-exact matches to a polynucleotide sequence of the invention (the 'Query'). A sequence from the UniGene database (the 'Subject') was said to be an exact match if it contained a segment of 50 nucleotides in length such that 48 of those nucleotides were in the same order as found in the Query sequence. If all of the matches that met this criteria were in the same UniGene cluster, and mapping data was available for this cluster, it is indicated in Table 1B under the heading "Cytologic Band". Where a cluster had been further localized to a distinct cytologic band, that band is disclosed; where no banding information was available, but the gene had been localized to a single chromosome, the chromosome is disclosed.

[88] Once a presumptive chromosomal location was determined for a polynucleotide of the invention, an associated disease locus was identified by comparison with a database of diseases which have been experimentally associated with genetic loci. The database used was the Morbid Map, derived from OMIM<sup>TM</sup> ("Online Mendelian Inheritance in Man"; McKusick-Nathans Institute for Genetic Medicine, Johns Hopkins University (Baltimore, MD) and National Center for Biotechnology Information, National Library of Medicine (Bethesda, MD) 2000; World Wide Web URL: http://www.ncbi.nlm.nih.gov/omim/). If the putative chromosomal location of a polynucleotide of the invention (Query sequence)

was associated with a disease in the Morbid Map database, an OMIM reference identification number was noted in column 10, Table 1B, labelled "OMIM Disease Reference(s). Table 5 is a key to the OMIM reference identification numbers (column 1), and provides a description of the associated disease in Column 2.

[89] Table 1C summarizes additional polynucleotides encompassed by the invention (including cDNA clones related to the sequences (Clone ID NO:Z), contig sequences (contig identifier (Contig ID:) contig nucleotide sequence identifiers (SEQ ID NO:X)), and genomic sequences (SEQ ID NO:B). The first column provides a unique clone identifier, "Clone ID NO:Z", for a cDNA clone related to each contig sequence. The second column provides the sequence identifier, "SEO ID NO:X", for each contig sequence. The third column provides a unique contig identifier, "Contig ID:" for each contig sequence. The fourth column, provides a BAC identifier "BAC ID NO:A" for the BAC clone referenced in the corresponding row of the table. The fifth column provides the nucleotide sequence identifier, "SEQ ID NO:B" for a fragment of the BAC clone identified in column four of the corresponding row of the table. The sixth column, "Exon From-To", provides the location (i.e., nucleotide position numbers) within the polynucleotide sequence of SEO ID NO:B which delineate certain polynucleotides of the invention that are also exemplary members of polynucleotide sequences that encode polypeptides of the invention (e.g., polypeptides containing amino acid sequences encoded by the polynucleotide sequences delineated in column six, and fragments and variants thereof).

## TABLE 1C

Clone ID	SEQ ID No:X	CONTIG ID	BAC ID: A	SEQ ID NO:B	EXON From-To
H6BSF56	11	762968	AC069362	1019	1-13
H6BSF56	11	762968	AC027584	1020	1-162
H6BSF56	11	762968	AC011101	1021	1-100
H6BSF56	11	762968	AC073446	1022	1-140
H6BSF56	11	762968	AC026556	1023	1-114
H6BSF56	11	762968	AL136171	1024	1-6
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HWDAG96	506	796743	AL356652	2149	1-638
					793-854
HWDAJ01	507	794016	AC015551	2150	1-670
HWDAJ01	507	794016	AC019214	2151	1-670
HWHPB78	508	740778	AL157945	2152	1-300
					364-790
					1344-1519
					1584-1709
	1				2403-2580
					4780-4968
					5485-5559
					5960-6128 6243-6955
					7258-7317
				+	9073-9145
					9404-9544
					10342-10513
					10746-11354
					12004-12578
					12863-13087
					13224-13382
					13993-14047
	İ				14319-14444
					14753-14878
					15465-15713
					16007-16123
					17413-17740
					17817-18127
					18231-18634
					18771-18881
					19945-20231
					21024-21169
		1			23112-23363
1					23692-24413
HWHPB78	508	740778	AC026283	2153	1-292
					353-776
					1340-1506
					1568-1696
					2408-2534
			]		4767-4955 5472-5546
					5957-6293
				Į	6373-7085
					7386-7445
					9201-9273
					9532-9672
					10470-10641
					10470-10041
					12131-12705
					12990-13214
					12770-13214

					13351-13509
					14119-14173
					14445-14570
					14879-15004
					15604-15844
					16133-16253
					17540-17867
					17944-18254
		}			18356-18755
					18892-19002
					20066-20352
					21146-21308
					23235-23486
		- 10-0	17.4570.45		23813-24533
HWHPB78	508	740778	AL157945	2154	1-490
HWHPB78	508	740778	AC026283	2155	1-318
HYABC84	509	789854	AL132825	2156	1-2512
	-				2604-2740
					2974-3241
HYABC84	509	789854	AL132825	2157	1-553
				·	1059-1263
	ļ ·				3121-3476
•					5284-5734
					6284-6513
					6786-7426
		]			8674-8733
					10656-10933
					11453-11555
					12991-13079
:					13839-14281
					14527-14827
					15156-15685
					15835-16046
					16166-16604 16736-19566
					19658-19794
					20028-20295
TIMADORA	509	789854	AL132825	2158	1-188
HYABC84	510	865064	AL132825	2159	1-2512
HYABC84	310	803004	AL132023	2139	2604-2740
					2974-3241
HVADC94	510	965064	AT 122925	2160	1-553
HYABC84	510	865064	AL132825	2100	1059-1263
					3121-3476
					5284-5734
					6284-6513
					6786-7426
					8674-8733
					10656-10933
					11453-11555
					12991-13079
					13839-14281
					14527-14827
	1				15156-15685
		ŀ			15835-16046
					16166-16604
					16736-19566
					19658-19794
					20028-20295
HYABC84	510	865064	AL132825	2161	1-188
LILLADU04	1510	003004	AL132023	12101	1-100

- [90] Tables 1D and 1E: The polynucleotides or polypeptides, or agonists or antagonists of the present invention can be used in assays to test for one or more biological activities. If these polynucleotides and polypeptides do exhibit activity in a particular assay, it is likely that these molecules may be involved in the diseases associated with the biological activity. Thus, the polynucleotides or polypeptides, or agonists or antagonists could be used to treat the associated disease.
- [91] The present invention encompasses methods of preventing, treating, diagnosing, or ameliorating a disease or disorder. In preferred embodiments, the present invention encompasses a method of treating a disease or disorder listed in the "Preferred Indications" columns of Table 1D and Table 1E; comprising administering to a patient in which such treatment, prevention, or amelioration is desired a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) in an amount effective to treat, prevent, diagnose, or ameliorate the disease or disorder. The first and second columns of Table 1D show the "Gene No." and "cDNA Clone ID No.", respectively, indicating certain nucleic acids and proteins (or antibodies against the same) of the invention (including polynucleotide, polypeptide, and antibody fragments or variants thereof) that may be used in preventing, treating, diagnosing, or ameliorating the disease(s) or disorder(s) indicated in the corresponding row in Column 3 of Table 1D.
- [92] In another embodiment, the present invention also encompasses methods of preventing, treating, diagnosing, or ameliorating a disease or disorder listed in the "Preferred Indications" column of Table 1D and Table 1E; comprising administering to a patient combinations of the proteins, nucleic acids, or antibodies of the invention (or fragments or variants thereof), sharing similar indications as shown in the corresponding rows in Column 3 of Table 1D.
- [93] The "Preferred Indications" columns of Table 1D and Table 1E describe diseases, disorders, and/or conditions that may be treated, prevented, diagnosed, or ameliorated by a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof).
- [94] The recitation of "Cancer" in the "Preferred Indications" columns indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof) may be used for example, to diagnose, treat, prevent, and/or

ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., leukemias, cancers, and/or as described below under "Hyperproliferative Disorders").

[95] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Cancer" recitation in the "Preferred Indication" column of Table 1D may be used for example, to diagnose, treat, prevent, and/or ameliorate a neoplasm located in a tissue selected from the group consisting of: colon, abdomen, bone, breast, digestive system, liver, pancreas, prostate, peritoneum, lung, blood (e.g., leukemia), endocrine glands (adrenal, parathyroid, pituitary, testicles, ovary, thymus, thyroid), uterus, eye, head and neck, nervous (central and peripheral), lymphatic system, pelvic, skin, soft tissue, spleen, thoracic, and urogenital.

[96] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Cancer" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a pre-neoplastic condition, selected from the group consisting of: hyperplasia (e.g., endometrial hyperplasia and/or as described in the section entitled "Hyperproliferative Disorders"), metaplasia (e.g., connective tissue metaplasia, atypical metaplasia, and/or as described in the section entitled "Hyperproliferative Disorders"), and/or dysplasia (e.g., cervical dysplasia, and bronchopulmonary dysplasia).

[97] In another specific embodiment, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Cancer" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a benign dysproliferative disorder selected from the group consisting of: benign tumors, fibrocystic conditions, tissue hypertrophy, and/or as described in the section entitled "Hyperproliferative Disorders".

[98] The recitation of "Immune/Hematopoietic" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders"), blood disorders (e.g., as described below under "Immune Activity" "Cardiovascular Disorders" and/or "Blood-Related Disorders"), and infections (e.g., as described below under "Infectious Disease").

[99] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having the "Immune/Hematopoietic" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat,

prevent, and/or ameliorate a disease or disorder selected from the group consisting of: anemia, pancytopenia, leukopenia, thrombocytopenia, leukemias, Hodgkin's disease, non-Hodgkin's lymphoma, acute lymphocytic anemia (ALL), plasmacytomas, multiple myeloma, Burkitt's lymphoma, arthritis, asthma, AIDS, autoimmune disease, rheumatoid arthritis, granulomatous disease, immune deficiency, inflammatory bowel disease, sepsis, neutropenia, neutrophilia, psoriasis, immune reactions to transplanted organs and tissues, systemic lupus erythematosis, hemophilia, hypercoagulation, diabetes mellitus, endocarditis, meningitis, Lyme Disease, and allergies.

[100] The recitation of "Reproductive" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders"), and disorders of the reproductive system (e.g., as described below under "Reproductive System Disorders").

[101] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Reproductive" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: cryptorchism, prostatitis, inguinal hernia, varicocele, leydig cell tumors, verrucous carcinoma, prostatitis, malacoplakia, Peyronie's disease, penile carcinoma, squamous cell hyperplasia, dysmenorrhea, ovarian adenocarcinoma, Turner's syndrome, mucopurulent cervicitis, Sertoli-leydig tumors, ovarian cancer, uterine cancer, pelvic inflammatory disease, testicular cancer, prostate cancer, Klinefelter's syndrome, Young's syndrome, premature ejaculation, diabetes mellitus, cystic fibrosis, Kartagener's syndrome, testicular atrophy, testicular feminization, anorchia, ectopic testis, epididymitis, orchitis, gonorrhea, syphilis, testicular torsion, vasitis nodosa, germ cell tumors, stromal tumors, dysmenorrhea, retroverted uterus, endometriosis, fibroids, adenomyosis, anovulatory bleeding, amenorrhea, Cushing's syndrome, hydatidiform moles, Asherman's syndrome, premature menopause, precocious puberty, uterine polyps, dysfunctional uterine bleeding, cervicitis, chronic cervicitis, mucopurulent cervicitis, cervical dysplasia, cervical polyps, Nabothian cysts, cervical erosion, cervical incompetence, cervical neoplasms, pseudohermaphroditism, and premenstrual syndrome.

[102] The recitation of "Musculoskeletal" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the

invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders"), and disorders of the immune system (e.g., as described below under "Immune Activity").

[103] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Musculoskeletal" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: bone cancers (e.g., osteochondromas, benign chondromas, chondroblastoma, chondromyxoid fibromas, osteoid osteomas, giant cell tumors, multiple myeloma, osteosarcomas), Paget's Disease, rheumatoid arthritis, systemic lupus erythematosus, osteomyelitis, Lyme Disease, gout, bursitis, tendonitis, osteoporosis, osteoarthritis, muscular dystrophy, mitochondrial myopathy, cachexia, and multiple sclerosis.

[104] The recitation of "Cardiovascular" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders"), and disorders of the cardiovascular system (e.g., as described below under "Cardiovascular Disorders").

[105] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Cardiovascular" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: myxomas, fibromas, rhabdomyomas, cardiovascular abnormalities (e.g., congenital heart defects, cerebral arteriovenous malformations, septal defects), heart disease (e.g., heart failure, congestive heart disease, arrhythmia, tachycardia, fibrillation, pericardial Disease, endocarditis), cardiac arrest, heart valve disease (e.g., stenosis, regurgitation, prolapse), vascular disease (e.g., hypertension, coronary artery disease, angina, aneurysm, arteriosclerosis, peripheral vascular disease), hyponatremia, hypernatremia, hypokalemia, and hyperkalemia.

[106] The recitation of "Mixed Fetal" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent,

and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders").

[107] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Mixed Fetal" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: spina bifida, hydranencephaly, neurofibromatosis, fetal alcohol syndrome, diabetes mellitus, PKU, Down's syndrome, Patau syndrome, Edwards syndrome, Turner syndrome, Apert syndrome, Carpenter syndrome, Conradi syndrome, Crouzon syndrome, cutis laxa, Cornelia de Lange syndrome, Ellis-van Creveld syndrome, Holt-Oram syndrome, Kartagener syndrome, Meckel-Gruber syndrome, Noonan syndrome, Pallister-Hall syndrome, Rubinstein-Taybi syndrome, Scimitar syndrome, Smith-Lemli-Opitz syndrome, thromocytopenia-absent radius (TAR) syndrome, Treacher Collins syndrome, Williams syndrome, Hirschsprung's disease, Meckel's diverticulum, polycystic kidney disease, Turner's syndrome, and gonadal dysgenesis, Klippel-Feil syndrome, Ostogenesis imperfecta, muscular dystrophy, Tay-Sachs disease, Wilm's tumor, neuroblastoma, and retinoblastoma.

[108] The recitation of "Excretory" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders") and renal disorders (e.g., as described below under "Renal Disorders").

[109] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Excretory" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: bladder cancer, prostate cancer, benign prostatic hyperplasia, bladder disorders (e.g., urinary incontinence, urinary retention, urinary obstruction, urinary tract Infections, interstitial cystitis, prostatitis, neurogenic bladder, hematuria), renal disorders (e.g., hydronephrosis, proteinuria, renal failure, pyelonephritis, urolithiasis, reflux nephropathy, and unilateral obstructive uropathy).

[110] The recitation of "Neural/Sensory" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the

invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders") and diseases or disorders of the nervous system (e.g., as described below under "Neural Activity and Neurological Diseases").

[111] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Neural/Sensory" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: brain cancer (e.g., brain stem glioma, brain tumors, central nervous system (Primary) lymphoma, central nervous system lymphoma, cerebellar astrocytoma, and cerebral astrocytoma, neurodegenerative disorders (e.g., Alzheimer's Disease, Creutzfeldt-Jakob Disease, Parkinson's Disease, and Idiopathic Presenile Dementia), encephalomyelitis, cerebral malaria, meningitis, metabolic brain diseases (e.g., phenylketonuria and pyruvate carboxylase deficiency), cerebellar ataxia, ataxia telangiectasia, and AIDS Dementia Complex, schizophrenia, attention deficit disorder, hyperactive attention deficit disorder, autism, and obsessive compulsive disorders.

[112] The recitation of "Respiratory" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders") and diseases or disorders of the respiratory system (e.g., as described below under "Respiratory Disorders").

[113] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Respiratory" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: cancers of the respiratory system such as larynx cancer, pharynx cancer, trachea cancer, epiglottis cancer, lung cancer, squamous cell carcinomas, small cell (oat cell) carcinomas, large cell carcinomas, and adenocarcinomas. Allergic reactions, cystic fibrosis, sarcoidosis, histiocytosis X, infiltrative lung diseases (e.g., pulmonary fibrosis and lymphoid interstitial pneumonia), obstructive airway diseases (e.g., asthma, emphysema, chronic or acute bronchitis), occupational lung diseases (e.g., silicosis and asbestosis), pneumonia, and pleurisy.

[114] The recitation of "Endocrine" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders") and diseases or disorders of the respiratory system (e.g., as described below under "Respiratory Disorders"), renal disorders (e.g., as described below under "Renal Disorders"), and disorders of the endocrine system (e.g., as described below under "Endocrine Disorders".

[115] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having an "Endocrine" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: cancers of endocrine tissues and organs (e.g., cancers of the hypothalamus, pituitary gland, thyroid gland, parathyroid glands, pancreas, adrenal glands, ovaries, and testes), diabetes (e.g., diabetes insipidus, type I and type II diabetes mellitus), obesity, disorders related to pituitary glands (e.g., hyperpituitarism, hypopituitarism, and pituitary dwarfism), hypothyroidism, hyperthyroidism, goiter, reproductive disorders (e.g. male and female infertility), disorders related to adrenal glands (e.g., Addison's Disease, corticosteroid deficiency, and Cushing's Syndrome), kidney cancer (e.g., hypernephroma, transitional cell cancer, and Wilm's tumor), diabetic nephropathy, interstitial nephritis, polycystic kidney disease, glomerulonephritis (e.g., IgM mesangial proliferative glomerulonephritis and glomerulonephritis caused by autoimmune disorders; such as Goodpasture's syndrome), and nephrocalcinosis.

[116] The recitation of "Digestive" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders") and diseases or disorders of the gastrointestinal system (e.g., as described below under "Gastrointestinal Disorders".

[117] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Digestive" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: ulcerative colitis, appendicitis, Crohn's disease, hepatitis, hepatic encephalopathy, portal hypertension,

cholelithiasis, cancer of the digestive system (e.g., biliary tract cancer, stomach cancer, colon cancer, gastric cancer, pancreatic cancer, cancer of the bile duct, tumors of the colon (e.g., polyps or cancers), and cirrhosis), pancreatitis, ulcerative disease, pyloric stenosis, gastroenteritis, gastritis, gastric atropy, benign tumors of the duodenum, distension, irritable bowel syndrome, malabsorption, congenital disorders of the small intestine, bacterial and parasitic infection, megacolon, Hirschsprung's disease, aganglionic megacolon, acquired megacolon, colitis, anorectal disorders (e.g., anal fistulas, hemorrhoids), congenital disorders of the liver (e.g., Wilson's disease, hemochromatosis, cystic fibrosis, biliary atresia, and alpha1-antitrypsin deficiency), portal hypertension, cholelithiasis, and jaundice.

[118] The recitation of "Connective/Epithelial" in the "Preferred Indication" column indicates that the corresponding nucleic acid and protein, or antibody against the same, of the invention (or fragment or variant thereof), may be used for example, to diagnose, treat, prevent, and/or ameliorate diseases and/or disorders relating to neoplastic diseases (e.g., as described below under "Hyperproliferative Disorders"), cellular and genetic abnormalities (e.g., as described below under "Diseases at the Cellular Level "), angiogenesis (e.g., as described below under "Anti-Angiogenesis Activity "), and or to promote or inhibit regeneration (e.g., as described below under "Regeneration "), and wound healing (e.g., as described below under "Wound Healing and Epithelial Cell Proliferation").

[119] In specific embodiments, a protein, nucleic acid, or antibody of the invention (or fragment or variant thereof) having a "Connective/Epithelial" recitation in the "Preferred Indication" column of Table 1D, may be used for example, to diagnose, treat, prevent, and/or ameliorate a disease or disorder selected from the group consisting of: connective tissue metaplasia, mixed connective tissue disease, focal epithelial hyperplasia, epithelial metaplasia, mucoepithelial dysplasia, graft v. host disease, polymyositis, cystic hyperplasia, cerebral dysplasia, tissue hypertrophy, Alzheimer's disease, lymphoproliferative disorder, Waldenstron's macroglobulinemia, Crohn's disease, pernicious anemia, idiopathic Addison's disease, glomerulonephritis, bullous pemphigoid, Sjogren's syndrome, diabetes mellitus, cystic fibrosis, osteoblastoma, osteoclastoma, osteosarcoma, chondrosarcoma, osteoporosis, osteocarthritis, periodontal disease, wound polychondritis, vasculitis, polyarteritis nodosa, Wegener's healing, relapsing granulomatosis, cellulitis, rheumatoid arthritis, psoriatic arthritis, discoid lupus erythematosus, systemic lupus erythematosus, scleroderma, CREST syndrome, Sjogren's syndrome, polymyositis, dermatomyositis, mixed connective tissue disease, relapsing polychondritis, vasculitis, Henoch-Schonlein syndrome, erythema nodosum, polyarteritis nodosa, temporal (giant cell) arteritis, Takayasu's arteritis, Wegener's granulomatosis, Reiter's syndrome, Behcet's syndrome, ankylosing spondylitis, cellulitis, keloids, Ehler Danlos syndrome, Marfan syndrome, pseudoxantoma elasticum, osteogenese imperfecta, chondrodysplasias, epidermolysis bullosa, Alport syndrome, and cutis laxa.

**TABLE 1D** 

Gene No.	Clone ID	Preferred Indications
1	H6BSF56	Cancer
2	H6EDM64	Cancer
3	H6EEC72	Cancer
4	HACAB68	Connective/Epithelial,
		Immune/Hematopoetic
5	HACBJ56	Cancer
6	HACBS22	Cancer
7	HADDE71	Cancer
8	HADDJ13	Connective/Epithelial
9	HADMB15	Cancer
10	HAGBQ12	Excretory,
		Neural/Sensory
11	HAGDW20	Neural/Sensory,
		Reproductive
12	HAGEG10	Cancer
13	HAGEQ79	Cancer
14	HAGFS57	Cancer
15	HAGHN57	Cancer
16	HAHEA15	Cardiovascular
17	HAJAA47	Immune/Hematopoetic
18	HAJAY92	Cancer
19	HAJBV67	Cancer
20	HAJCH70	Cancer
21	HAOAG15	Cancer
22	HAQAI92	Digestive,
	,	Mixed Fetal,
		Reproductive
23	HAQCE11	Reproductive
24	HATBI94	Cancer
25	HATCB45	Endocrine,
		Immune/Hematopoetic
26	HATCD80	Endocrine,
		Reproductive
27	HATCI03	Endocrine,
		Immune/Hematopoetic,
		Neural/Sensory
28	HATEH20	Cancer
29	HBAGD86	Cancer
30	HBCJL35	Cancer
31	HBDAB91	Digestive,
		Immune/Hematopoetic
32	HBDAB91	Digestive,

Immune/Hematopoetic			
			Immune/Hematopoetic
35			
According to the content of the co			
Neural/Sensory   Santa   HBIAC29   Cancer   Cancer   Digestive,	35		
	36	HBHAA81	
BIAC29			
Big   Big	37		Cancer
Immune/Hematopoetic, Neural/Sensory	38		Cancer
Neural/Sensory	39	HBICW51	
HBJAB02   Cancer			
HBJAC65			
HBJBM12	40		
HBJCR46	41	HBJAC65	
HBIDS79	42	HBJBM12	Immune/Hematopoetic
HBJDW56	43	HBJCR46	Cancer
HBJEL16	44	HBJDS79	
47         HBJFK45         Immune/Hematopoetic           48         HBJG20         Cancer           49         HBJKD16         Cancer           50         HBMBM96         Cancer           51         HBMBX01         Cancer           52         HBMTM11         Cancer           53         HBMTX26         Immune/Hematopoetic           54         HBMTY48         Immune/Hematopoetic           54         HBMTY48         Immune/Hematopoetic           55         HBMUT44         Cardiovascular, Immune/Hematopoetic, Reproductive           55         HBMWE61         Immune/Hematopoetic           56         HBNAX40         Cancer           58         HBNBJR76         Cancer           59         HBQAB79         Neural/Sensory           60         HBQAC57         Neural/Sensory           61         HBSAK32         Cancer           62         HBXCM66         Cardiovascular, Neural/Sensory, Reproductive           63         HBXCX15         Immune/Hematopoetic, Neural/Sensory           64         HCDDL48         Musculoskeletal           65         HCE2H52         Immune/Hematopoetic, Neural/Sensory, Reproductive           68	45	HBJDW56	Immune/Hematopoetic
HBJIG20	46	HBJEL16	Cancer
HBJKD16	47	HBJFK45	Immune/Hematopoetic
SO	48	HBJIG20	Cancer
51         HBMBX01         Cancer           52         HBMTM11         Cancer           53         HBMTX26         Immune/Hematopoetic           54         HBMTY48         Immune/Hematopoetic,           54         HBMTY48         Immune/Hematopoetic,           55         HBMUH74         Cardiovascular,           55         HBMUH74         Cardiovascular,           56         HBMWE61         Immune/Hematopoetic           57         HBNAX40         Cancer           59         HBNAX40         Cancer           60         HBQAB79         Neural/Sensory           61         HBACK32         Cancer           62         HBXCM66         Cardiovascular,           Neural/Sensory,         Neural/Sensory           Reproductive         Neural/Sensory           64         HCDCY76         Cancer           65         HCDDL48         Musculoskeletal           66         HCE1G78         Cancer           67         HCE2H52         Immune/Hematopoetic,           Neural/Sensory,         Reproductive           68         HCE3B04         Cancer           69         HCE5F78         Immune/Hematopoetic,	49	HBJKD16	Cancer
51         HBMBX01         Cancer           52         HBMTM11         Cancer           53         HBMTX26         Immune/Hematopoetic           54         HBMTY48         Immune/Hematopoetic,           54         HBMTY48         Immune/Hematopoetic,           55         HBMUH74         Cardiovascular,           55         HBMUH74         Cardiovascular,           56         HBMWE61         Immune/Hematopoetic           57         HBNAX40         Cancer           59         HBNAX40         Cancer           60         HBQAB79         Neural/Sensory           61         HBACK32         Cancer           62         HBXCM66         Cardiovascular,           Neural/Sensory,         Neural/Sensory           Reproductive         Neural/Sensory           64         HCDCY76         Cancer           65         HCDDL48         Musculoskeletal           66         HCE1G78         Cancer           67         HCE2H52         Immune/Hematopoetic,           Neural/Sensory,         Reproductive           68         HCE3B04         Cancer           69         HCE5F78         Immune/Hematopoetic,	50	HBMBM96	Cancer
52         HBMTM11         Cancer           53         HBMTX26         Immune/Hematopoetic           54         HBMTY48         Immune/Hematopoetic, Reproductive           55         HBMUH74         Cardiovascular, Immune/Hematopoetic, Reproductive           56         HBMWE61         Immune/Hematopoetic           57         HBNAX40         Cancer           58         HBNBJ76         Cancer           59         HBQAB79         Neural/Sensory           60         HBQAC57         Neural/Sensory           61         HBSAK32         Cancer           62         HBXCM66         Cardiovascular, Neural/Sensory, Reproductive           63         HBXCX15         Immune/Hematopoetic, Neural/Sensory           64         HCDCY76         Cancer           65         HCDDL48         Musculoskeletal           66         HCE1G78         Cancer           67         HCE2H52         Immune/Hematopoetic, Neural/Sensory, Reproductive           68         HCE5F78         Immune/Hematopoetic, Neural/Sensory           70         HCEDR26         Digestive, Immune/Hematopoetic, Neural/Sensory           71         HCEEC25         Mixed Fetal, Neural/Sensory		HBMBX01	Cancer
53         HBMTX26         Immune/Hematopoetic           54         HBMTY48         Immune/Hematopoetic, Reproductive           55         HBMUH74         Cardiovascular, Immune/Hematopoetic, Reproductive           56         HBMWE61         Immune/Hematopoetic           57         HBNAX40         Cancer           58         HBNBJ76         Cancer           59         HBQAB79         Neural/Sensory           60         HBQAC57         Neural/Sensory           61         HBSAK32         Cancer           62         HBXCM66         Cardiovascular, Neural/Sensory, Reproductive           63         HBXCX15         Immune/Hematopoetic, Neural/Sensory           64         HCDCY76         Cancer           65         HCDDL48         Musculoskeletal           66         HCE1G78         Cancer           67         HCE2H52         Immune/Hematopoetic, Neural/Sensory, Reproductive           68         HCE3B04         Cancer           69         HCE5F78         Immune/Hematopoetic, Neural/Sensory           70         HCEDR26         Digestive, Immune/Hematopoetic, Neural/Sensory           71         HCEEC25         Mixed Fetal, Neural/Sensory			Cancer
54         HBMTY48         Immune/Hematopoetic, Reproductive           55         HBMUH74         Cardiovascular, Immune/Hematopoetic, Reproductive           56         HBMWE61         Immune/Hematopoetic           57         HBNAX40         Cancer           58         HBNBJ76         Cancer           59         HBQAB79         Neural/Sensory           60         HBQAC57         Neural/Sensory           61         HBSAK32         Cancer           62         HBXCM66         Cardiovascular, Neural/Sensory, Reproductive           63         HBXCX15         Immune/Hematopoetic, Neural/Sensory           64         HCDCY76         Cancer           65         HCDDL48         Musculoskeletal           66         HCE1G78         Cancer           67         HCE2H52         Immune/Hematopoetic, Neural/Sensory, Reproductive           68         HCE3B04         Cancer           69         HCE5F78         Immune/Hematopoetic, Neural/Sensory           70         HCEDR26         Digestive, Immune/Hematopoetic, Neural/Sensory           71         HCEECQ25         Mixed Fetal, Neural/Sensory			
Reproductive		HBMTY48	
S5			
Immune/Hematopoetic, Reproductive	55	HBMUH74	
56         HBMWE61         Immune/Hematopoetic           57         HBNAX40         Cancer           58         HBNBJ76         Cancer           59         HBQAB79         Neural/Sensory           60         HBQAC57         Neural/Sensory           61         HBSAK32         Cancer           62         HBXCM66         Cardiovascular, Neural/Sensory, Reproductive           63         HBXCX15         Immune/Hematopoetic, Neural/Sensory           64         HCDCY76         Cancer           65         HCDDL48         Musculoskeletal           66         HCE1G78         Cancer           67         HCE2H52         Immune/Hematopoetic, Neural/Sensory, Reproductive           68         HCE3B04         Cancer           69         HCE5F78         Immune/Hematopoetic, Neural/Sensory           70         HCEDR26         Digestive, Immune/Hematopoetic, Neural/Sensory           71         HCEEQ25         Mixed Fetal, Neural/Sensory			Immune/Hematopoetic,
57         HBNAX40         Cancer           58         HBNBJ76         Cancer           59         HBQAB79         Neural/Sensory           60         HBQAC57         Neural/Sensory           61         HBSAK32         Cancer           62         HBXCM66         Cardiovascular, Neural/Sensory, Reproductive           63         HBXCX15         Immune/Hematopoetic, Neural/Sensory           64         HCDCY76         Cancer           65         HCDDL48         Musculoskeletal           66         HCE1G78         Cancer           67         HCE2H52         Immune/Hematopoetic, Neural/Sensory, Reproductive           68         HCE3B04         Cancer           69         HCE5F78         Immune/Hematopoetic, Neural/Sensory           70         HCEDR26         Digestive, Immune/Hematopoetic, Neural/Sensory           71         HCEEQ25         Mixed Fetal, Neural/Sensory			Reproductive
58         HBNBJ76         Cancer           59         HBQAB79         Neural/Sensory           60         HBQAC57         Neural/Sensory           61         HBSAK32         Cancer           62         HBXCM66         Cardiovascular, Neural/Sensory, Reproductive           63         HBXCX15         Immune/Hematopoetic, Neural/Sensory           64         HCDCY76         Cancer           65         HCDDL48         Musculoskeletal           66         HCE1G78         Cancer           67         HCE2H52         Immune/Hematopoetic, Neural/Sensory, Reproductive           68         HCE3B04         Cancer           69         HCE5F78         Immune/Hematopoetic, Neural/Sensory           70         HCEDR26         Digestive, Immune/Hematopoetic, Neural/Sensory           71         HCEEC79         Neural/Sensory           72         HCEEQ25         Mixed Fetal, Neural/Sensory	56	HBMWE61	Immune/Hematopoetic
59         HBQAB79         Neural/Sensory           60         HBQAC57         Neural/Sensory           61         HBSAK32         Cancer           62         HBXCM66         Cardiovascular, Neural/Sensory, Reproductive           63         HBXCX15         Immune/Hematopoetic, Neural/Sensory           64         HCDCY76         Cancer           65         HCDDL48         Musculoskeletal           66         HCE1G78         Cancer           67         HCE2H52         Immune/Hematopoetic, Neural/Sensory, Reproductive           68         HCE3B04         Cancer           69         HCE5F78         Immune/Hematopoetic, Neural/Sensory           70         HCEDR26         Digestive, Immune/Hematopoetic, Neural/Sensory           71         HCEEC25         Mixed Fetal, Neural/Sensory	57	HBNAX40	Cancer
60         HBQAC57         Neural/Sensory           61         HBSAK32         Cancer           62         HBXCM66         Cardiovascular, Neural/Sensory, Reproductive           63         HBXCX15         Immune/Hematopoetic, Neural/Sensory           64         HCDCY76         Cancer           65         HCDDL48         Musculoskeletal           66         HCE1G78         Cancer           67         HCE2H52         Immune/Hematopoetic, Neural/Sensory, Reproductive           68         HCE3B04         Cancer           69         HCE5F78         Immune/Hematopoetic, Neural/Sensory           70         HCEDR26         Digestive, Immune/Hematopoetic, Neural/Sensory           71         HCEEQ25         Mixed Fetal, Neural/Sensory	58	HBNBJ76	Cancer
60         HBQAC57         Neural/Sensory           61         HBSAK32         Cancer           62         HBXCM66         Cardiovascular, Neural/Sensory, Reproductive           63         HBXCX15         Immune/Hematopoetic, Neural/Sensory           64         HCDCY76         Cancer           65         HCDDL48         Musculoskeletal           66         HCE1G78         Cancer           67         HCE2H52         Immune/Hematopoetic, Neural/Sensory, Reproductive           68         HCE3B04         Cancer           69         HCE5F78         Immune/Hematopoetic, Neural/Sensory           70         HCEDR26         Digestive, Immune/Hematopoetic, Neural/Sensory           71         HCEEQ25         Mixed Fetal, Neural/Sensory	59	HBQAB79	Neural/Sensory
61         HBSAK32         Cancer           62         HBXCM66         Cardiovascular, Neural/Sensory, Reproductive           63         HBXCX15         Immune/Hematopoetic, Neural/Sensory           64         HCDCY76         Cancer           65         HCDDL48         Musculoskeletal           66         HCE1G78         Cancer           67         HCE2H52         Immune/Hematopoetic, Neural/Sensory, Reproductive           68         HCE3B04         Cancer           69         HCE5F78         Immune/Hematopoetic, Neural/Sensory           70         HCEDR26         Digestive, Immune/Hematopoetic, Neural/Sensory           71         HCEEC9         Neural/Sensory           72         HCEQ25         Mixed Fetal, Neural/Sensory		HBQAC57	Neural/Sensory
62         HBXCM66         Cardiovascular, Neural/Sensory, Reproductive           63         HBXCX15         Immune/Hematopoetic, Neural/Sensory           64         HCDCY76         Cancer           65         HCDDL48         Musculoskeletal           66         HCE1G78         Cancer           67         HCE2H52         Immune/Hematopoetic, Neural/Sensory, Reproductive           68         HCE3B04         Cancer           69         HCE5F78         Immune/Hematopoetic, Neural/Sensory           70         HCEDR26         Digestive, Immune/Hematopoetic, Neural/Sensory           71         HCEEC79         Neural/Sensory           72         HCEEQ25         Mixed Fetal, Neural/Sensory	61		Cancer
Reproductive		HBXCM66	Cardiovascular,
63         HBXCX15         Immune/Hematopoetic, Neural/Sensory           64         HCDCY76         Cancer           65         HCDDL48         Musculoskeletal           66         HCE1G78         Cancer           67         HCE2H52         Immune/Hematopoetic, Neural/Sensory, Reproductive           68         HCE3B04         Cancer           69         HCE5F78         Immune/Hematopoetic, Neural/Sensory           70         HCEDR26         Digestive, Immune/Hematopoetic, Neural/Sensory           71         HCEEC79         Neural/Sensory           72         HCEEQ25         Mixed Fetal, Neural/Sensory			Neural/Sensory,
Neural/Sensory			Reproductive
64         HCDCY76         Cancer           65         HCDDL48         Musculoskeletal           66         HCE1G78         Cancer           67         HCE2H52         Immune/Hematopoetic,	63	HBXCX15	Immune/Hematopoetic,
65         HCDDL48         Musculoskeletal           66         HCE1G78         Cancer           67         HCE2H52         Immune/Hematopoetic, Neural/Sensory, Reproductive           68         HCE3B04         Cancer           69         HCE5F78         Immune/Hematopoetic, Neural/Sensory           70         HCEDR26         Digestive, Immune/Hematopoetic, Neural/Sensory           71         HCEEC79         Neural/Sensory           72         HCEEQ25         Mixed Fetal, Neural/Sensory			Neural/Sensory
65         HCDDL48         Musculoskeletal           66         HCE1G78         Cancer           67         HCE2H52         Immune/Hematopoetic, Neural/Sensory, Reproductive           68         HCE3B04         Cancer           69         HCE5F78         Immune/Hematopoetic, Neural/Sensory           70         HCEDR26         Digestive, Immune/Hematopoetic, Neural/Sensory           71         HCEEC79         Neural/Sensory           72         HCEEQ25         Mixed Fetal, Neural/Sensory	64	HCDCY76	
66         HCE1G78         Cancer           67         HCE2H52         Immune/Hematopoetic, Neural/Sensory, Reproductive           68         HCE3B04         Cancer           69         HCE5F78         Immune/Hematopoetic, Neural/Sensory           70         HCEDR26         Digestive, Immune/Hematopoetic, Neural/Sensory           71         HCEEC79         Neural/Sensory           72         HCEEQ25         Mixed Fetal, Neural/Sensory			Musculoskeletal
67 HCE2H52 Immune/Hematopoetic, Neural/Sensory, Reproductive  68 HCE3B04 Cancer  69 HCE5F78 Immune/Hematopoetic, Neural/Sensory  70 HCEDR26 Digestive, Immune/Hematopoetic, Neural/Sensory  71 HCEEC79 Neural/Sensory  72 HCEEQ25 Mixed Fetal, Neural/Sensory			Cancer
Neural/Sensory, Reproductive  68			
68 HCE3B04 Cancer  69 HCE5F78 Immune/Hematopoetic, Neural/Sensory  70 HCEDR26 Digestive, Immune/Hematopoetic, Neural/Sensory  71 HCEEC79 Neural/Sensory  72 HCEEQ25 Mixed Fetal, Neural/Sensory			Neural/Sensory,
69 HCE5F78 Immune/Hematopoetic, Neural/Sensory  70 HCEDR26 Digestive, Immune/Hematopoetic, Neural/Sensory  71 HCEEE79 Neural/Sensory  72 HCEEQ25 Mixed Fetal, Neural/Sensory			Reproductive
Neural/Sensory  Neural/Sensory  Digestive, Immune/Hematopoetic, Neural/Sensory  HCEE79 Neural/Sensory  HCEEQ25 Mixed Fetal, Neural/Sensory	68	HCE3B04	
70 HCEDR26 Digestive, Immune/Hematopoetic, Neural/Sensory  71 HCEEE79 Neural/Sensory  72 HCEEQ25 Mixed Fetal, Neural/Sensory	69	HCE5F78	
Immune/Hematopoetic, Neural/Sensory  71 HCEEE79 Neural/Sensory  72 HCEEQ25 Mixed Fetal, Neural/Sensory			
Neural/Sensory  71 HCEEE79 Neural/Sensory  72 HCEEQ25 Mixed Fetal, Neural/Sensory	70	HCEDR26	
71 HCEE79 Neural/Sensory 72 HCEEQ25 Mixed Fetal, Neural/Sensory			
72 HCEEQ25 Mixed Fetal, Neural/Sensory			
Neural/Sensory	71	HCEEE79	
Neural/Sensory	72	HCEEQ25	
73 HCEEU18 Cancer			Neural/Sensory
	73	HCEEU18	Cancer

	Livennes	
74	HCEFZ82	Cancer
75	HCEGX05	Cancer
76	HCFLN88	Cancer
77	HCFLT90	Cancer
78	HCHAB84	Cancer
79	HCMSX51	Cancer
80	HCNCO11	Digestive
81	HCNSD29	Cardiovascular,
		Digestive,
		Immune/Hematopoetic
82	HCQBH72	Digestive,
		Excretory,
		Immune/Hematopoetic
83	HCQCC96	Cancer
84	HCQCJ56	Cancer
85	HCQCM24	Cancer
86	HCRAY10	Cancer
87	HCRBF72	Cancer
88	HCRNF78	Cancer
89	HCUAF85	Immune/Hematopoetic
90	HCUCF89	Immune/Hematopoetic
91	HCUCK44	Cancer
92	HCUDD64	Cancer
93	HCWAE64	Immune/Hematopoetic
94	HCWFU39	Endocrine,
) <del>74</del>	IICWI 039	Immune/Hematopoetic,
		Neural/Sensory
95	HCWUL09	Immune/Hematopoetic,
93	HCW OLU9	Neural/Sensory
96	HDHAA42	Cancer
97	HDHEB76	Cancer
98	HDPCW16	Cancer
99	HDPDI72	
99	HDPD1/2	Digestive, Immune/Hematopoetic
100	HDPDJ58	Cancer
100		the state of the s
101	HDPFF10	Cancer
102	HDPFU43	Cancer
103	HDPFY18	Cancer
104	HDPGE24	Cancer
105	HDPIU94	Cancer
106	HDPOC24	Cancer
107	HDPOL37	Immune/Hematopoetic,
		Reproductive
108	HDPOO76	Cancer
109	HDPPD93	Cancer
110	HDPPQ30	Immune/Hematopoetic
111	HDPPW82	Immune/Hematopoetic
112	HDPXN20	Immune/Hematopoetic
113	HDQHM36	Immune/Hematopoetic
114	HDTAU35	Immune/Hematopoetic
115	HDTAV54	Cancer
116	HDTFX18	Immune/Hematopoetic,
		Reproductive
117	HDTGW48	Immune/Hematopoetic,
		Reproductive
118	HDTLM18	Immune/Hematopoetic
119	HE2CA60	Cancer
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	T	
120	HE2CA60	Cancer
121	HE2CH58	Digestive,
100	115001430	Mixed Fetal
122	HE2CM39	Cancer
123 124	HE2HC60	Cancer
125	HE2PO93	Cancer Mixed Fetal
126	HE6AU52	
127	HE6CS65 HE6DO92	Cancer
12/	HE0DO92	Immune/Hematopoetic, Mixed Fetal
128	HE6EY13	Cancer
129	HE6FU11	Mixed Fetal,
12)	11201 011	Neural/Sensory,
		Respiratory
130	HE6FV29	Cancer
131	HE8FC45	Cancer
132	HE8FC45	Cancer
133	HE8FD92	Cancer
134	HE8FD92	Cancer
135	HE8FD92	Cancer
136	HE8FD92	Cancer
137	HE8FD92	Cancer
138	HE8SG96	Mixed Fetal,
		Musculoskeletal
139	HE8TY46	Cancer
140	HE9CY05	Mixed Fetal
141	HE9EA10	Cancer
142	HE9GG20	Cancer
143	HEBCI18	Cancer
144	HEBCY54	Cancer
145	HEBDF77	Neural/Sensory
146	HEBDQ91	Neural/Sensory
147	HEBFR46	Cancer
148	HEBGE07	Neural/Sensory
149	HEGAU15	Excretory,
		Immune/Hematopoetic,
150	7777 4 770 5	Reproductive
150	HELAT35	Cardiovascular,
151	TIEL DITEA	Mixed Fetal
152	HELBU54 HELGG84	Cardiovascular Cancer
153	HELGG84	Cancer
154	HEMEY47	Cardiovascular
155	HEOMC46	Immune/Hematopoetic
156	HEPBA14	Reproductive
157	HEQAH80	Cancer
158	HEQBF89	Reproductive
159	HETCI16	Cancer
160	HETDW58	Cancer
161	HETEY67	Connective/Epithelial,
	IIII   III	Reproductive
162	HFCDW95	Cancer
163	HFCEI04	Neural/Sensory
164	HFCFD04	Cancer
165	HFCFE20	Cancer
166	HFEAY59	Connective/Epithelial
167	HFGAJ16	Cancer

168	HFIHZ75	Cancer
169	HFIJA29	Cancer
170	HFIJA68	Cancer
171	HFKES05	Cancer
172	HFKEU12	Excretory
173	HFPCZ55	Cancer
174	HFPDR62	Immune/Hematopoetic,
		Neural/Sensory
175	HFPDS07	Cancer
176	HFRAB10	Excretory,
		Immune/Hematopoetic,
		Neural/Sensory
177	HFTBM38	Cancer
178	HFTDH56	Cancer
179	HFVGK35	Cancer
180	HFVHW43	Digestive
181	HFXAV37	Immune/Hematopoetic,
		Neural/Sensory
182	HFXBN86	Neural/Sensory
183	HFXBT66	Neural/Sensory
184	HFXFZ46	Neural/Sensory
185	HGBER72	Cancer
186	HGBEY14	Cancer
187	HGBGN34	Connective/Epithelial,
107	11GDGI134	Digestive,
		Reproductive
188	HGBHP91	Digestive
189	HGCAC19	Cancer
190	HGCAC19	Cancer
191	HGCAC19	Cancer
192	HHEAK45	Cancer
193		
194	HHEGS55	Immune/Hematopoetic
	HHEOW19	Cancer
195	HHFFF87	Cancer
196	HHFFL34	Cancer
197	HHFFS40	Cancer
198	HHGCS78	Immune/Hematopoetic
199	HHGDT26	Immune/Hematopoetic,
		Reproductive
200	HHPFU28	Cancer
201	HHPSA85	Cancer
202	HHSBI06	Cancer
203	HHSBI65	Cancer
204	HHSDI53	Cancer
205	HHSFC09	Cancer
206	HHSGL28	Cancer
207	HILCA24	Digestive,
		Immune/Hematopoetic,
		Reproductive
208	HILCA24	Digestive,
		Immune/Hematopoetic,
		Reproductive
209	HISAT67	Cancer
210	HJBCU75	Cancer
211	HJMAA03	Cancer
212	НЈМАV41	Cancer
213	НЈМАҮ90	Cancer
	1	

214	HJPBE39	Cancer
215	НЈРВК28	Cancer
216	НЈРСН08	Cancer
217	HKABU43	Cancer
218	HKACI79	Cancer
219	HKAFF50	Cancer
220	HKGBF25	Cancer
221	HKIXC44	Cancer
222	HKMLK03	Digestive,
		Excretory,
		Immune/Hematopoetic
223	HKMLM95	Cancer
224	HKTAB41	Digestive,
		Excretory
225	HLDBG17	Cancer
226	HLDCA54	Cancer
227	HLDQU79	Cancer
228	HLDRT09	Cancer
229	HLHAP05	Immune/Hematopoetic,
	1	Neural/Sensory,
		Respiratory
230	HLHCS23	Respiratory
231	HLIBO72	Digestive
232	HLICE88	Digestive,
	<u> </u>	Mixed Fetal
233	HLICO10	Cancer
234	HLJBS28	Cancer
235	HLMBW89	Cancer
236	HLMGP50	Digestive, Immune/Hematopoetic
237	НЬМЈВ64	Cancer
238	HLMMX62	Immune/Hematopoetic,
		Neural/Sensory,
1	:	Reproductive
1		
239	HLQAS12	Cancer
240	HLQCL64	Cancer
240 241	HLQCL64 HLQCX36	Cancer Digestive
240	HLQCL64	Cancer Digestive Digestive,
240 241	HLQCL64 HLQCX36	Cancer Digestive Digestive, Immune/Hematopoetic,
240 241 242	HLQCL64 HLQCX36 HLWAF06	Cancer Digestive Digestive, Immune/Hematopoetic, Reproductive
240 241 242 243	HLQCL64 HLQCX36 HLWAF06 HLWAU42	Cancer Digestive Digestive, Immune/Hematopoetic, Reproductive Cancer
240 241 242 243 244	HLQCL64 HLQCX36 HLWAF06 HLWAU42 HLWAU42	Cancer Digestive Digestive, Immune/Hematopoetic, Reproductive Cancer Cancer
240 241 242 243 244 245	HLQCL64 HLQCX36 HLWAF06 HLWAU42 HLWAU42 HLWAV47	Cancer Digestive Digestive, Immune/Hematopoetic, Reproductive Cancer Cancer Cancer
240 241 242 243 244 245 246	HLQCL64 HLQCX36 HLWAF06 HLWAU42 HLWAU42 HLWAV47 HLWBB73	Cancer Digestive Digestive, Immune/Hematopoetic, Reproductive Cancer Cancer Cancer Cancer
240 241 242 243 244 245 246 247	HLQCL64 HLQCX36 HLWAF06  HLWAU42 HLWAU42 HLWAV47 HLWBB73 HLWCN37	Cancer Digestive Digestive, Immune/Hematopoetic, Reproductive Cancer Cancer Cancer Cancer Cancer Cancer
240 241 242 243 244 245 246 247 248	HLQCL64 HLQCX36 HLWAF06 HLWAU42 HLWAU42 HLWAV47 HLWBB73 HLWCN37 HLWDB73	Cancer Digestive Digestive, Immune/Hematopoetic, Reproductive Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer
240 241 242 243 244 245 246 247 248 249	HLQCL64 HLQCX36 HLWAF06  HLWAU42 HLWAU42 HLWAV47 HLWBB73 HLWCN37 HLWDB73 HLYDF73	Cancer Digestive Digestive, Immune/Hematopoetic, Reproductive Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Immune/Hematopoetic
240 241 242 243 244 245 246 247 248 249 250	HLQCL64 HLQCX36 HLWAF06  HLWAU42 HLWAU42 HLWAV47 HLWBB73 HLWCN37 HLWDB73 HLYDF73 HLYDF73	Cancer Digestive Digestive, Immune/Hematopoetic, Reproductive Cancer Cancer Cancer Cancer Cancer Immune/Hematopoetic Immune/Hematopoetic Immune/Hematopoetic
240 241 242 243 244 245 246 247 248 249 250 251	HLQCL64 HLQCX36 HLWAF06  HLWAU42 HLWAU42 HLWAV47 HLWBB73 HLWCN37 HLWDB73 HLYDF73 HLYEU59 HLYGB19	Cancer Digestive Digestive, Immune/Hematopoetic, Reproductive Cancer Cancer Cancer Cancer Cancer Immune/Hematopoetic Immune/Hematopoetic Immune/Hematopoetic Cancer
240 241 242 243 244 245 246 247 248 249 250 251 252	HLQCL64 HLQCX36 HLWAF06  HLWAU42 HLWAU42 HLWAV47 HLWBB73 HLWCN37 HLWDB73 HLYDF73 HLYDF73 HLYEU59 HLYGB19 HLYGE16	Cancer Digestive Digestive, Immune/Hematopoetic, Reproductive Cancer Cancer Cancer Cancer Cancer Immune/Hematopoetic Immune/Hematopoetic Immune/Hematopoetic Cancer Cancer
240 241 242 243 244 245 246 247 248 249 250 251 252 253	HLQCL64 HLQCX36 HLWAF06  HLWAU42 HLWAU42 HLWAV47 HLWBB73 HLWCN37 HLWDB73 HLYDF73 HLYEU59 HLYGB19 HLYGE16 HLYGY91	Cancer Digestive Digestive, Immune/Hematopoetic, Reproductive Cancer Immune/Hematopoetic Cancer Cancer Cancer
240 241 242  243 244 245 246 247 248 249 250 251 252 253 254	HLQCL64 HLQCX36 HLWAF06  HLWAU42 HLWAU42 HLWAV47 HLWBB73 HLWCN37 HLWDB73 HLYDF73 HLYEU59 HLYGB19 HLYGE16 HLYGY91 HMCAZ04	Cancer Digestive Digestive, Immune/Hematopoetic, Reproductive Cancer
240 241 242 243 244 245 246 247 248 249 250 251 252 253 254 255	HLQCL64 HLQCX36 HLWAF06  HLWAU42 HLWAU42 HLWAV47 HLWBB73 HLWCN37 HLWDB73 HLYDF73 HLYGB19 HLYGB19 HLYGB16 HLYGY91 HMCAZ04	Cancer Digestive Digestive, Immune/Hematopoetic, Reproductive Cancer Immune/Hematopoetic Immune/Hematopoetic Cancer Cancer Cancer Cancer Cancer
240 241 242 243 244 245 246 247 248 249 250 251 252 253 254 255 256	HLQCL64 HLQCX36 HLWAF06  HLWAU42 HLWAU42 HLWAV47 HLWBB73 HLWCN37 HLWDB73 HLYDF73 HLYEU59 HLYGB19 HLYGE16 HLYGY91 HMCAZ04 HMCAZ04	Cancer Digestive Digestive, Immune/Hematopoetic, Reproductive Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Immune/Hematopoetic Cancer Cancer Cancer Cancer Cancer Cancer Cancer
240 241 242  243 244 245 246 247 248 249 250 251 252 253 254 255 256 257	HLQCL64 HLQCX36 HLWAF06  HLWAU42 HLWAU42 HLWAV47 HLWBB73 HLWCN37 HLWDB73 HLYDF73 HLYGE16 HLYGE16 HLYGY91 HMCAZ04 HMCAZ04 HMCAZ04	Cancer Digestive Digestive, Immune/Hematopoetic, Reproductive Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Immune/Hematopoetic Immune/Hematopoetic Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer
240 241 242 243 244 245 246 247 248 249 250 251 252 253 254 255 256	HLQCL64 HLQCX36 HLWAF06  HLWAU42 HLWAU42 HLWAV47 HLWBB73 HLWCN37 HLWDB73 HLYDF73 HLYEU59 HLYGB19 HLYGE16 HLYGY91 HMCAZ04 HMCAZ04	Cancer Digestive Digestive, Immune/Hematopoetic, Reproductive Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Immune/Hematopoetic Cancer Cancer Cancer Cancer Cancer Cancer Cancer

	T	
260	HMDAB29	Digestive,
261	TIME A DATA	Neural/Sensory
261	HMDAD44	Connective/Epithelial,
		Immune/Hematopoetic, Neural/Sensory
262	ID (EDD02	Cancer
262	HMEBB82	
263	HMEDE24	Cancer
264	HMEDI90	Cancer
265	HMELM75	Cancer
266	HMIAK10	Neural/Sensory
267	HMIBF07	Neural/Sensory
268	HMICI80	Cardiovascular,
		Endocrine,
260	III (ICD(5	Neural/Sensory
269	HMICP65	Cancer
270	HMJAK70	Neural/Sensory
271	HMSBE04	Immune/Hematopoetic
272	HMSCL38	Digestive,
		Immune/Hematopoetic,
272		Neural/Sensory
273	HMSCR69	Cancer
274	HMSHC86	Immune/Hematopoetic
275	HMSHU20	Immune/Hematopoetic,
276	TIN COLUMN S	Reproductive
276	HMSHY25	Immune/Hematopoetic
277	HMTAB77	Cancer
278	HMUAE26	Cancer
279	HMUAN45	Cancer
280	HMVBC31	Cancer
281	HMVDU15	Cancer
282	HMWBL03	Cancer
283	HMWJF53	Cancer
284	HNEAK81	Immune/Hematopoetic
285	HNECL22	Cancer
286	HNECW49	Immune/Hematopoetic
287	HNEDH88	Immune/Hematopoetic
288	HNFAC50	Cancer
289	HNFGR08	Immune/Hematopoetic
290	HNFHF34	Cancer
291	HNGAK51	Immune/Hematopoetic
292	HNGAM58	Immune/Hematopoetic
293	HNGBH53	Immune/Hematopoetic
294	HNGDQ38	Immune/Hematopoetic
295	HNGDX18	Cancer
296	HNGDY34	Immune/Hematopoetic
297	HNGEA34	Digestive,
		Immune/Hematopoetic
298	HNGEQ75	Immune/Hematopoetic,
		Neural/Sensory
299	HNGGA68	Immune/Hematopoetic,
		Musculoskeletal
300	HNGGP65	Immune/Hematopoetic
301	HNGHZ69	Immune/Hematopoetic
302	HNGIV64	Immune/Hematopoetic
303	HNGJB41	Immune/Hematopoetic
304	HNGKT41	Immune/Hematopoetic
305	HNGMW45	Immune/Hematopoetic

306	HNGNK44	Immune/Hematopoetic
307	HNGNO53	Immune/Hematopoetic
308	HNGPJ25	Immune/Hematopoetic,
		Mixed Fetal,
		Musculoskeletal
309	HNHEN82	Immune/Hematopoetic
310	HNHFE71	Immune/Hematopoetic
311	HNHGK22	Immune/Hematopoetic
312	HNHHB10	Immune/Hematopoetic,
		Reproductive
313	HNHKS19	Immune/Hematopoetic,
		Reproductive
314	HNTBT17	Cancer
315	HNTMH79	Cancer
316	HOABP31	Cancer
317	HOABP31	Cancer
318	HOACG07	Cancer
319	HODAG07	Reproductive
320	HODBB70	Reproductive
321	HODBV05	Cancer
322	HODCZ32	Reproductive
323	HOEBK60	Cancer
324	HOFAA78	Reproductive
325	HOFNB74	Reproductive
326	HOFNU55	Reproductive
<b></b>		Reproductive
327	HOGBF01	Cancer
328	HORBS82	
329	HORBV76	Cardiovascular,
		Immune/Hematopoetic, Reproductive
220	HOSDO75	Cancer
330		4.45
331	HOSEC25	Immune/Hematopoetic, Musculoskeletal,
		Reproductive
222	HOCEIGI	Digestive,
332	HOSEI81	Musculoskeletal
333	HOSEJ94	Cancer
334	HOUCA21	Connective/Epithelial,
334	HOUCAZI	Immune/Hematopoetic.
		Illinune/Hematopoetic,
		Musculoskeletal
225	HOUDEON	Musculoskeletal
335	HOUDE92	Cancer
336	HOUDR07	Cancer Cancer
336 337	HOUDR07 HOUED72	Cancer Cancer Connective/Epithelial
336 337 338	HOUDR07 HOUED72 HOUFS04	Cancer Cancer Connective/Epithelial Cancer
336 337 338 339	HOUDR07 HOUED72 HOUFS04 HOUHI25	Cancer Cancer Connective/Epithelial Cancer Cancer
336 337 338	HOUDR07 HOUED72 HOUFS04	Cancer Cancer Connective/Epithelial Cancer Cancer Musculoskeletal,
336 337 338 339 340	HOUDR07 HOUED72 HOUFS04 HOUHI25 HOVBD85	Cancer Cancer Connective/Epithelial Cancer Cancer Musculoskeletal, Reproductive
336 337 338 339	HOUDR07 HOUED72 HOUFS04 HOUHI25	Cancer Cancer Connective/Epithelial Cancer Cancer Musculoskeletal, Reproductive Immune/Hematopoetic,
336 337 338 339 340	HOUDR07 HOUED72 HOUFS04 HOUHI25 HOVBD85	Cancer Cancer Connective/Epithelial Cancer Cancer Musculoskeletal, Reproductive Immune/Hematopoetic, Reproductive
336 337 338 339 340 341	HOUDR07 HOUED72 HOUFS04 HOUHI25 HOVBD85 HPCAB41 HPCAL26	Cancer Cancer Connective/Epithelial Cancer Cancer Musculoskeletal, Reproductive Immune/Hematopoetic, Reproductive Cancer
336 337 338 339 340 341 342 343	HOUDR07 HOUED72 HOUFS04 HOUHI25 HOVBD85 HPCAB41 HPCAL26 HPEAD23	Cancer Cancer Connective/Epithelial Cancer Cancer Musculoskeletal, Reproductive Immune/Hematopoetic, Reproductive Cancer Cancer Cancer
336 337 338 339 340 341 342 343 344	HOUDR07 HOUED72 HOUFS04 HOUHI25 HOVBD85 HPCAB41 HPCAL26 HPEAD23 HPFBA54	Cancer Cancer Connective/Epithelial Cancer Cancer Musculoskeletal, Reproductive Immune/Hematopoetic, Reproductive Cancer Cancer Reproductive
336 337 338 339 340 341 342 343 344 345	HOUDR07 HOUED72 HOUFS04 HOUHI25 HOVBD85 HPCAB41 HPCAL26 HPEAD23 HPFBA54 HPFCI36	Cancer Cancer Connective/Epithelial Cancer Cancer Musculoskeletal, Reproductive Immune/Hematopoetic, Reproductive Cancer Cancer Reproductive Cancer Cancer
336 337 338 339 340 341 342 343 344 345 346	HOUDR07 HOUED72 HOUFS04 HOUHI25 HOVBD85 HPCAB41 HPCAL26 HPEAD23 HPFBA54 HPFCI36 HPFDI37	Cancer Cancer Connective/Epithelial Cancer Cancer Musculoskeletal, Reproductive Immune/Hematopoetic, Reproductive Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer
336 337 338 339 340 341 342 343 344 345 346 347	HOUDR07 HOUED72 HOUFS04 HOUHI25 HOVBD85 HPCAB41 HPCAL26 HPEAD23 HPFBA54 HPFCI36 HPFDI37 HPIAA80	Cancer Cancer Connective/Epithelial Cancer Cancer Musculoskeletal, Reproductive Immune/Hematopoetic, Reproductive Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer
336 337 338 339 340 341 342 343 344 345 346	HOUDR07 HOUED72 HOUFS04 HOUHI25 HOVBD85 HPCAB41 HPCAL26 HPEAD23 HPFBA54 HPFCI36 HPFDI37	Cancer Cancer Connective/Epithelial Cancer Cancer Musculoskeletal, Reproductive Immune/Hematopoetic, Reproductive Cancer Cancer Cancer Cancer Cancer Cancer Cancer Cancer

250	Tiprovic	
350	HPJBU43	Reproductive
351	HPJCW58	Reproductive
352	HPMBX22	Cancer
353	HPMCJ84	Reproductive
354	HPMCV30	Cancer
355	HPMFH77	Cancer
356	HPQAX38	Cardiovascular
357	HPQAX38	Cardiovascular
358	HPQCB83	Cancer
359	HPQCC53	Cancer
360	HPRBH85	Cancer
361	HPRCA64	Cancer
362	HPRCD35	Cancer
363	HPTRM02	Cancer
364	HPWBA29	Reproductive
365	HPWDK06	Cancer
366	HRAAD30	Cancer
367	HRADA42	Cancer
368	HRADF49	Cancer
369	HRADN25	Cancer
370	HRADT25	Digestive,
		Excretory
371	HRDAI17	Cancer
372	HRDDQ39	Cancer
373	HRDER22	Cancer
374	HRDEX93	Cancer
375	HRDFK37	Cancer
376	HRGBD54	Cancer
377	HROEA08	Cancer
378	HSAVA08	Immune/Hematopoetic
379	HSAVW42	Cancer
380	HSAWN53	Immune/Hematopoetic
381	HSAWZ40	Immune/Hematopoetic
382	HSAYC41	Excretory,
		Immune/Hematopoetic,
	<u> </u>	Reproductive
383	HSDZM54	Cancer
384	HSHBF76	Cancer
385	HSIFG47	Digestive
386	HSJBY32	Immune/Hematopoetic,
		Musculoskeletal,
		Neural/Sensory
387	HSKDR27	Cancer
388	HSLHG78	Cancer
389	HSLHX15	Musculoskeletal
390	HSNAP85	Cancer
391	HSNAZ09	Cancer
392	HSNBM34	Digestive
393	HSOAH16	Digestive
394	HSQBF66	Cancer
395	HSQDO85	Cancer
396	HSQES57	Cancer
397	HSRBE06	Cancer
398	HSSDI26	Musculoskeletal
399	HSSEA64	Cancer
400	HSSEF77	Cancer
401	HSSFE38	Cancer
<u> </u>	1	L. T. T. T. T. T. T. T. T. T. T. T. T. T.

402	HSSGJ58	Musculoskeletal
403	HSWBE76	Cancer
404	HSXCP38	
404	полсто	Cardiovascular, Neural/Sensory
405	HSYBI06	Cancer
406	HT1SC27	Digestive,
400	11113027	Immune/Hematopoetic,
		Reproductive
407	HT3BF49	Immune/Hematopoetic
408	HT4FV41	Cancer
409	HT5FX79	Cancer
410	HT5GR59	Cancer
411	HTAEI78	Immune/Hematopoetic
412	HTDAA78	Cancer
413	HTEAG62	Digestive,
		Immune/Hematopoetic,
		Reproductive
414	HTECB02	Cancer
415	HTECC15	Cancer
416	HTEDF18	Reproductive
417	HTEDJ28	Cancer
418	HTEDS12	Cardiovascular,
		Immune/Hematopoetic,
		Reproductive
419	HTEED26	Cancer
420	HTEED26	Cancer
421	HTEEF26	Cancer
422	HTEEF26	Cancer
423	HTEEW69	Reproductive
424	HTEGS07	Reproductive
425	HTEGS11	Cancer
426	HTEHA56	Cancer
427	HTEHU59	Cancer
428	HTEJD29	Reproductive
429	HTEKM46	Cancer
430	HTEMQ17	Cancer
431	HTENR63	Cancer
432	HTGGM44	Immune/Hematopoetic,
		Musculoskeletal
433	HTHBZ06	Cancer
434	HTLAP64	Cancer
435	HTLBT80	Cancer
436	HTLDA84	Reproductive
437	HTLDN29	Cancer
438	HTLDU78	Reproductive
439	HTLEC82	Cancer
440	HTLEM16	Cancer
441	HTLEV48	Reproductive
442	HTLFA13	Musculoskeletal,
442	1101 5150	Reproductive
443	HTLFI73	Cancer
444	HTLGI89	Cancer
445	HTLIF11	Cancer
446	HTLIF12	Excretory,
447	HTI IE12	Reproductive
<del></del>	HTLIF12	Excretory, Reproductive
		Reproductive

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448	HTLIF12	Excretory,
	1707 1512	Reproductive
449	HTLIF12	Excretory,
		Reproductive
450	HTLIF12	Excretory,
		Reproductive
451	HTLIF12	Excretory,
		Reproductive
452	HTNAM63	Endocrine
453	HTNBK13	Cancer
454	HTOAI50	Immune/Hematopoetic
455	HTOAM11	Immune/Hematopoetic,
		Neural/Sensory
456	HTODH57	Immune/Hematopoetic
457	HTODH83	Immune/Hematopoetic
458	HTOEV16	Cancer
459	HTOGR38	Immune/Hematopoetic
460	HTOHO21	Immune/Hematopoetic
461	HTOHQ05	Immune/Hematopoetic
462	HTOJL95	Cancer
463	HTOJL95	Cancer
464	HTPDU17	Cancer
465	HTSFJ32	Immune/Hematopoetic
466	HTTCB60	Cancer
467	HTTEE41	Cancer
468	HTTEZ02	Cancer
469	HTWEH94	Immune/Hematopoetic
470	HTXBD09	Cancer
471	HTXDB22	Cancer
472	HTXDC38	Cancer
473	HTXDC77	Cancer
474	HTXDD61	Cancer
475	HTXDG92	Cancer
476	HTXET11	Immune/Hematopoetic
477	HTXFA72	Immune/Hematopoetic
478	HTXJY08	Cancer
479	HTXKF95	Cancer
480	HTXMZ07	Cancer
481	HUFCL31	Digestive,
401	HOPCLSI	Immune/Hematopoetic
482	HUKBT67	Cancer
483	HUKDF20	Cardiovascular,
403	HUKDI 20	Reproductive
484	HUKDY82	Cancer
485	HUSCJ14	
		Cancer
486	HUSGL67	Cancer
487	HUSGU40	Cancer
488	HUSIR18	Cancer
489	HUVDJ48	Digestive,
400	TIME A TIO	Reproductive
490	HWAAI12	Cancer
491	HWBBQ70	Immune/Hematopoetic,
102	I I I I I I I I I I I I I I I I I I I	Neural/Sensory
492	HWBCN36	Immune/Hematopoetic
493	HWBDJ08	Cancer
494	HWBFX16	Immune/Hematopoetic
495	HWDAC26	Connective/Epithelial,

		Immune/Hematopoetic, Neural/Sensory	
496	HWDAG96	Cancer	
497	HWDAJ01	Connective/Epithelial	
498	HWHPB78	Cancer	
499	HYABC84	Cancer	
500	HYABC84	Cancer	

[120] Table 1E provides information related to biological activities and preferred indications for polynucleotides and polypeptides of the invention (including antibodies, agonists, and/or antagonists thereof). Table 1E also provides information related to assays which may be used to test polynucleotides and polypeptides of the invention (including antibodies, agonists, and/or antagonists thereof) for the corresponding biological activities. The first column ("Gene No.") provides the gene number in the application for each clone identifier. The second column ("cDNA Clone ID No:Z") provides the unique clone identifier for each clone as previously described and indicated in Tables 1A, 1B, 1C, and 1D. The third column ("AA SEQ ID NO:Y") indicates the Sequence Listing SEQ ID Number for polypeptide sequences encoded by the corresponding cDNA clones (also as indicated in Tables 1A, 1B, and 2). The fourth column ("Biological Activity") indicates a biological activity corresponding to the indicated polypeptides (or polynucleotides encoding said polypeptides). The fifth column ("Exemplary Activity Assay") further describes the corresponding biological activity and also provides information pertaining to the various types of assays which may be performed to test, demonstrate, or quantify the corresponding biological activity. The sixth column ("Preferred Indictions") describes particular embodiments of the invention as well as indications (e.g. pathologies, diseases, disorders, abnormalities, etc.) for which polynucleotides and polypeptides of the invention (including antibodies, agonists, and/or antagonists thereof) may be used in detecting, diagnosing, preventing, and/or treating.

[121] Table 1E describes the use of, inter alia, FMAT technology for testing or demonstrating various biological activities. Fluorometric microvolume assay technology (FMAT) is a fluorescence-based system which provides a means to perform nonradioactive cell- and bead-based assays to detect activation of cell signal transduction pathways. This technology was designed specifically for ligand binding and immunological assays. Using this technology, fluorescent cells or beads at the bottom of the well are detected as localized areas of concentrated fluorescence using a data processing system. Unbound flurophore comprising the background signal is ignored,

allowing for a wide variety of homogeneous assays. FMAT technology may be used for peptide ligand binding assays, immunofluorescence, apoptosis, cytotoxicity, and bead-based immunocapture assays. See, Miraglia S et. al., "Homogeneous cell and bead based assays for highthroughput screening using flourometric microvolume assay technology," Journal of Biomolecular Screening; 4:193-204 (1999). In particular, FMAT technology may be used to test, confirm, and/or identify the ability of polypeptides (including polypeptide fragments and variants) to activate signal transduction pathways. For example, FMAT technology may be used to test, confirm, and/or identify the ability of polypeptides to upregulate production of immunomodulatory proteins (such as, for example, interleukins, GM-CSF, Rantes, and Tumor Necrosis factors, as well as other cellular regulators (e.g. insulin)).

[122] Table 1E also describes the use of kinase assays for testing, demonstrating, or quantifying biological activity. In this regard, the phosphorylation and de-phosphorylation of specific amino acid residues (e.g. Tyrosine, Serine, Threonine) on cell-signal transduction proteins provides a fast, reversible means for activation and de-activation of cellular signal transduction pathways. Moreover, cell signal transduction via phosphorylation/de-phosphorylation is crucial to the regulation of a wide variety of cellular processes (e.g. proliferation, differentiation, migration, apoptosis, etc.). Accordingly, kinase assays provide a powerful tool useful for testing, confirming, and/or identifying polypeptides (including polypeptide fragments and variants) that mediate cell signal transduction events via protein phosphorylation. See e.g., Forrer, P., Tamaskovic R., and Jaussi, R. "Enzyme-Linked Immunosorbent Assay for Measurement of JNK, ERK, and p38 Kinase Activities" Biol. Chem. 379(8-9): 1101-1110 (1998).

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## TARLE 1E

Gene	cDNA Clone ID	AA SEQ ID	Biological Activity	Exemplary Activity Assay	Preferred Indications
1	H6BSF56	515	Activation of	Assays for the activation of transcription	A preferred embodiment of the invention includes a
			transcription through	through the Serum Response Element	method for inhibiting (e.g., reducing) TNF alpha
			serum response element	(SRE) are well-known in the art and may	production. An alternative preferred embodiment of the
			in immune cells (such	be used or routinely modified to assess the	invention includes a method for stimulating (e.g.,
			as T-cells).	ability of polypeptides of the invention	increasing) TNF alpha production. Preferred indications
				(including antibodies and agonists or	include blood disorders (e.g., as described below under
				antagonists of the invention) to regulate	"Immune Activity", "Blood-Related Disorders", and/or
				the serum response factors and modulate	"Cardiovascular Disorders"), Highly preferred indications
				the expression of genes involved in	include autoimmune diseases (e.g., rheumatoid arthritis,
				growth. Exemplary assays for	systemic lupus erythematosis, Crohn's disease, multiple
				transcription through the SRE that may be	sclerosis and/or as described below), immunodeficiencies
				used or routinely modified to test SRE	(e.g., as described below), boosting a T cell-mediated
				activity of the polypeptides of the	immune response, and suppressing a T cell-mediated
				invention (including antibodies and	immune response. Additional highly preferred indications
				agonists or antagonists of the invention)	include inflammation and inflammatory disorders, and
				include assays disclosed in Berger et al.,	treating joint damage in patients with rheumatoid arthritis.
				Gene 66:1-10 (1998); Cullen and Malm,	An additional highly preferred indication is sepsis.
				Methods in Enzymol 216:362-368 (1992);	Highly preferred indications include neoplastic diseases
				Henthorn et al., Proc Natl Acad Sci USA	(e.g., leukemia, lymphoma, and/or as described below
				85:6342-6346 (1988); and Black et al.,	under "Hyperproliferative Disorders"). Additionally,
				Virus Genes 12(2):105-117 (1997), the	highly preferred indications include neoplasms and
	,			content of each of which are herein	cancers, such as, for example, leukemia, lymphoma,
				incorporated by reference in its entirety. T	melanoma, glioma (e.g., malignant glioma), solid tumors,
				cells that may be used according to these	and prostate, breast, lung, colon, pancreatic, esophageal,
				assays are publicly available (e.g., through	stomach, brain, liver and urinary cancer. Other preferred
				the ATCC). Exemplary mouse T cells that	indications include benign dysproliferative disorders and
				may be used according to these assays	pre-neoplastic conditions, such as, for example,
				include the CTLL cell line, which is an IL-	hyperplasia, metaplasia, and/or dysplasia. Preferred
				2 dependent suspension culture of T cells	indications include anemia, pancytopenia, leukopenia,
				with cytotoxic activity.	thrombocytopenia, Hodgkin's disease, acute lymphocytic

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					anemia (ALL), plasmacytomas, multiple myeloma, Burkitt's lymphoma, arthritis, AIDS, granulomatous disease, inflammatory bowel disease, neutropenia, neutrophilia, psoriasis, suppression of immune reactions to transplanted organs and tissues, hemophilia, hypercoagulation, diabetes mellitus, endocarditis, meningitis, Lyme Disease, cardiac reperfusion injury, and asthma and allergy. An additional preferred indication is infection (e.g., an infectious disease as described below
2	H6EDM64	516	Insulin Secretion	Assays for measuring secretion of insulin are well-known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of	under "Infectious Disease").  A highly preferred indication is diabetes mellitus. An additional highly preferred indication is a complication associated with diabetes (e.g., diabetic retinopathy, diabetic nephropathy, kidney disease (e.g., renal failure, nephropathy and/or other diseases and disorders as
				the invention) to stimulate insulin secretion. For example, insulin secretion is measured by FMAT using anti-rat insulin antibodies. Insulin secretion from pancreatic beta cells is upregulated by glucose and also by certain proteins/peptides, and disregulation is a key component in diabetes. Exemplary	described in the Kenal Disorders section below), diabetic neuropathy, nerve disease and nerve damage (e.g., due to diabetic neuropathy), blood vessel blockage, heart disease, stroke, impotence (e.g., due to diabetic neuropathy or blood vessel blockage), seizures, mental confusion, drowsiness, nonketotic hyperglycemic-hyperosmolar coma, cardiovascular disease (e.g., heart disease, atherosclerosis, microvascular disease, hypertension,
				assays that may be used or routinely modified to test for stimulation of insulin secretion (from pancreatic cells) by polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include assays disclosed in: Shimizu, H., et al., Endocr J, 47(3):261-9 (2000); Salapatek, A.M., et al., Mol	stroke, and other diseases and disorders as described in the "Cardiovascular Disorders" section below), dyslipidemia, endocrine disorders (as described in the "Endocrine Disorders" section below), neuropathy, vision impairment (e.g., diabetic retinopathy and blindness), ulcers and impaired wound healing, and infection (e.g., infectious diseases and disorders as described in the "Infectious Diseases" section below, especially of the urinary tract and
				Endocrinol, 13(8):1305-17 (1999); Filipsson, K., et al., Ann N Y Acad Sci, 865:441-4 (1998); Olson, L.K., et al., J Biol Chem, 271(28):16544-52 (1996); and, Miraglia S et. al., Journal of Biomolecular Screening, 4:193-204 (1999), the contents	skin), carpal tunnel syndrome and Dupuytren's contracture). An additional highly preferred indication is obesity and/or complications associated with obesity. Additional highly preferred indications include weight loss or alternatively, weight gain. Aditional highly preferred indications are complications associated

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with insulin resistance.	A preferred embodiment of the invention includes a method for inhibiting (e.g., reducing) TNF alpha production. An alternative highly preferred embodiment of the invention includes a method for stimulating (e.g., increasing) TNF alpha production. Preferred indications include blood disorders (e.g., as described below under "Immune Activity", "Blood-Related Disorders", and/or "Cardiovascular Disorders"), Highly preferred indications include autoimmune diseases (e.g., rheumatoid arthritis, systemic lupus erythematosis, Crohn's disease, multiple systemic lupus erythematosis, Crohn's disease, multiple sclerosis and/or as described below), immunodeficiencies sclerosis and/or as described below), immunodeficiencies immune response. Additional highly preferred indications include inflammation and inflammatory disorders, and include inflammation and inflammatory disorders, and treating joint damage in patients with rheumatoid arthritis. An additional highly preferred indications include neoplastic diseases
of each of which is herein incorporated by reference in its entirety. Pancreatic cells that may be used according to these assays are publicly available (e.g., through the ATCC) and/or may be routinely generated. Exemplary pancreatic cells that may be used according to these assays include HITT15 Cells. HITT15 are an adherent epithelial cell line established from Syrian hamster islet cells transformed with SV40. These cells express glucagon, somatostatin, and glucocorticoid receptors. The cells secrete insulin, which is stimulated by glucose and glucagon and suppressed by somatostatin or glucocorticoids. ATTC# CRL-1777 Refs: Lord and Ashcroft. Biochem. J. 219: 847-551; Santerre et al. Proc. Natl. Acad.	Sci. USA 18: 4337-4343, 1701.  Assays for the activation of transcription through the Serum Response Element (SRE) are well-known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) to regulate serum response factors and modulate the expression of genes involved in growth and upregulate the function of growth-related genes in many cell types.  Exemplary assays for transcription through the SRE that may be used or routinely modified to test SRE activity of the polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include assays disclosed in the invention) include assays disclosed in
	Activation of transcription through serum response element in immune cells (such as natural killer cells).
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(e.g., leukemia, lymphoma, and/or as described below under "Hyperproliferative Disorders"). Additionally, highly preferred indications include neoplasms and cancers, such as, for example, leukemia, lymphoma, melanoma, glioma (e.g., malignant glioma), solid tumors, and prostate, breast, lung, colon, pancreatic, esophageal, stomach, brain, liver and urinary cancer. Other preferred indications include benign dysproliferative disorders and pre-neoplastic conditions, such as, for example, hyperplasia, metaplasia, and/or dysplasia. Preferred indications include anemia, pancytopenia, leukopenia, thrombocytopenia, Hodgkin's disease, acute lymphocytic anemia (ALL), plasmacytomas, multiple myeloma, Burkitt's lymphoma, arthritis, AIDS, granulomatous disease, inflammatory bowel disease, neutropenia, neutrophilia, psoriasis, suppression of immune reactions to transplanted organs and tissues, hemophilia, hypercoagulation, diabetes mellitus, endocarditis, meningitis, Lyme Disease, cardiac reperfusion injury, and asthma and allergy. An additional preferred indication is infection (e.g., an infectious disease as described below under "Infectious Disease").	Preferred indications include blood disorders (e.g., as described below under "Immune Activity", "Blood-Related Disorders", and/or "Cardiovascular Disorders"), and infection (e.g., an infectious disease as described below under "Infectious Disease"). Preferred indications include autoimmune diseases (e.g., rheumatoid arthritis, systemic lupus erythematosis, multiple sclerosis and/or as described below), immunodeficiencies (e.g., as described below), boosting a T cell-mediated immune response, and suppressing a T cell-mediated immune response. Additional preferred indications include inflammatory disorders. Highly preferred indications and/or as described below under "Hyperproliferative Disorders"). Highly preferred
and Malm, Methods in Enzymol 216:362-368 (1992); Henthorn et al., Proc Natl Acad Sci USA 85:6342-6346 (1988); Benson et al., J Immunol 153(9):3862-3873 (1994); and Black et al., Virus Genes 12(2):105-117 (1997), the content of each of which are herein incorporated by reference in its entirety. T cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary T cells that may be used according to these assays include the NK-YT cell line, which is a human natural killer cell line with cytolytic and cytotoxic activity.	Assays for the activation of transcription through the cAMP response element are well-known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) to increase cAMP and regulate CREB transcription factors, and modulate expression of genes involved in a wide variety of cell functions. Exemplary assays for transcription through the cAMP response element that may be used or routinely modified to test cAMP-response element activity of polypeptides of the invention (including antibodies and
	Activation of transcription through cAMP response element in immune cells (such as T-cells).
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indications include neoplasms and cancers, such as, for example, leukemia, lymphoma (e.g., T cell lymphoma, Burkitt's lymphoma, non-Hodgkins lymphoma, Hodgkin's disease), melanoma, and prostate, breast, lung, colon, pancreatic, esophageal, stomach, brain, liver and urinary cancer. Other preferred indications include benign dysproliferative disorders and pre-neoplastic conditions, such as, for example, hyperplasia, metaplasia, and/or dysplasia. Preferred indications include anemia, pancytopenia, leukopenia, thrombocytopenia, acute lymphocytic anemia (ALL.), plasmacytomas, multiple myeloma, arthritis, AIDS, granulomatous disease, inflammatory bowel disease, sepsis, neutropenia, neutrophilia, psoriasis, suppression of immune reactions to transplanted organs and tissues, hemophilia, hypercoagulation, diabetes mellitus, endocarditis, meningitis, Lyme Disease, and asthma and allergy.	A preferred embodiment of the invention includes a method for inhibiting (e.g., reducing) TNF alpha production. An alternative preferred embodiment of the invention includes a method for stimulating (e.g., increasing) TNF alpha production. Preferred indications include blood disorders (e.g., as described below under "Tamune Activity", "Blood-Related Disorders", and/or "Cardiovascular Disorders"), Highly preferred indications include autoimmune diseases (e.g., rheumatoid arthritis, systemic lupus erythematosis, Crohn's disease, multiple sclerosis and/or as described below), immunodeficiencies (e.g., as described below), boosting a T cell-mediated immune response. Additional highly preferred indications include inflammation and inflammatory disorders, and treating joint damage in patients with rheumatoid arthritis. An additional highly preferred indication is sepsis. Highly preferred indications include neoplastic diseases (e.g., leukemia, lymphoma, and/or as described below
agonists or antagonists of the invention) include assays disclosed in Berger et al., Gene 66:1-10 (1998); Cullen and Malm, Methods in Enzymol 216:362-368 (1992); Henthorn et al., Proc Natl Acad Sci USA 85:6342-6346 (1988); Black et al., Virus Genes 15(2):105-117 (1997); and Belkowski et al., J Immunol 161(2):659-665 (1998), the contents of each of which are herein incorporated by reference in its entirety. T cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary mouse T cells that may be used according to these assays include the CTLL cell line, which is a suspension culture of IL-2 dependent cytotoxic T cells.	Assays for the activation of transcription through the Serum Response Element (SRE) are well-known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) to regulate the serum response factors and modulate the serum response factors and modulate the expression of genes involved in growth. Exemplary assays for transcription through the SRE that may be used or routinely modified to test SRE activity of the polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include assays disclosed in Berger et al., Gene 66:1-10 (1998); Cullen and Malm, Methods in Enzymol 216:362-368 (1992); Henthorn et al., Proc Natl Acad Sci USA
	Activation of transcription through serum response element in immune cells (such as T-cells).
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			-	85:6342-6346 (1988); and Black et al., Virus Genes 12(2):105-117 (1997), the content of each of which are herein incorporated by reference in its entirety. T cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary mouse T cells that may be used according to these assays include the CTLL cell line, which is an IL-2 dependent suspension culture of T cells with cytotoxic activity.	under "Hyperproliferative Disorders"). Additionally, highly preferred indications include neoplasms and cancers, such as, for example, leukemia, lymphoma, melanoma, glioma (e.g., malignant glioma), solid tumors, and prostate, breast, lung, colon, pancreatic, esophageal, stomach, brain, liver and urinary cancer. Other preferred indications include benign dysproliferative disorders and pre-neoplastic conditions, such as, for example, hyperplasia, metaplasia, and/or dysplasia. Preferred indications include anemia, pancytopenia, leukopenia, thrombocytopenia, Hodgkin's disease, acute lymphocytic anemia (ALL), plasmacytomas, multiple myeloma, Burkitt's lymphoma, arthritis, AIDS, granulomatous disease, inflammatory bowel disease, neutropenia, neutrophilia, psoriasis, suppression of immune reactions to transplanted organs and tissues, hemophilia, mortipis, Lyme Disease, cardiac reperfusion injury, and asthma and allergy. An additional preferred indication is infection (e.g., an infectious disease as described below under "Infectious Disease").
4	HACAB68	518	Activation of Endothelial Cell p38 or JNK Signaling Pathway.	Kinase assay. JNK and p38 kinase assays for signal transduction that regulate cell proliferation, activation, or apoptosis are well known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) to promote or inhibit cell proliferation, activation, and apoptosis. Exemplary assays for JNK and p38 kinase activity that may be used or routinely modified to test JNK and p38 kinase induced activity of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include the assays disclosed in Forrer et	A highly preferred embodiment of the invention includes a method for stimulating endothelial cell growth. An alternative highly preferred embodiment of the invention includes a method for inhibiting endothelial cell growth. A highly preferred embodiment of the invention includes a method for stimulating endothelial cell proliferation. An alternative highly preferred embodiment of the invention includes a method for inhibiting endothelial cell proliferation. A highly preferred embodiment of the invention includes a method for stimulating apoptosis of endothelial cells. An alternative highly preferred embodiment of the invention includes a method for inhibiting (e.g., decreasing) apoptosis of endothelial cells. A highly preferred embodiment of the invention includes a method for stimulating (e.g., increasing) endothelial cell activation.

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al. Biol Chem 379(8-9):1101-1110	An alternative highly preferred embodiment of the
(1998); Gupta et al., Exp Cell Res 247(2):	invention includes a method for inhibiting (e.g.,
495-504 (1999); Kyriakis JM, Biochem	decreasing) the activation of and/or inactivating
Soc Symp 64:29-48 (1999); Chang and	endothelial cells. A highly preferred embodiment of
Karin, Nature 410(6824):37-40 (2001);	the invention includes a method for stimulating
and Cobb MH, Prog Biophys Mol Biol	angiogenisis. An alternative highly preferred embodiment
71(3-4):479-500 (1999); the contents of	of the invention includes a method for inhibiting
each of which are herein incorporated by	angiogenesis. A highly preferred embodiment of the
 reference in its entirety. Endothelial cells	invention includes a method for reducing cardiac
that may be used according to these assays	hypertrophy. An alternative highly preferred embodiment
are publicly available (e.g., through the	of the invention includes a method for inducing cardiac
ATCC). Exemplary endothelial cells that	hypertrophy. Highly preferred indications include
may be used according to these assays	neoplastic diseases (e.g., as described below under
 include human umbilical vein endothelial	"Hyperproliferative Disorders"), and disorders of the
 cells (HUVEC), which are endothelial	cardiovascular system (e.g., heart disease, congestive heart
 cells which line venous blood vessels, and	failure, hypertension, aortic stenosis, cardiomyopathy,
are involved in functions that include, but	valvular regurgitation, left ventricular dysfunction,
are not limited to, angiogenesis, vascular	atherosclerosis and atherosclerotic vascular disease,
permeability, vascular tone, and immune	diabetic nephropathy, intracardiac shunt, cardiac
cell extravasation.	hypertrophy, myocardial infarction, chronic hemodynamic
	overload, and/or as described below under
	"Cardiovascular Disorders"). Highly preferred indications
	include cardiovascular, endothelial and/or angiogenic
	disorders (e.g., systemic disorders that affect vessels such
	as diabetes mellitus, as well as diseases of the vessels
	themselves, such as of the arteries, capillaries, veins and/or
	lymphatics). Highly preferred are indications that
	stimulate angiogenesis and/or cardiovascularization.
	Highly preferred are indications that inhibit angiogenesis
	and/or cardiovascularization. Highly preferred
	indications include antiangiogenic activity to treat solid
	tumors, leukemias, and Kaposi's sarcoma, and retinal
	disorders. Highly preferred indications include neoplasms
	and cancer, such as, Kaposi's sarcoma, hemangioma
	(capillary and cavernous), glomus tumors, telangiectasia,
	bacillary angiomatosis, hemangioendothelioma,
	angiosarcoma, haemangiopericytoma, lymphangioma,

lymphangiosarcoma. Highly preferred indications also include cancers such as, prostate, breast, lung, colon, pancreatic, esophageal, stomach, brain, liver, and urinary cancer. Preferred indications include benign dysproliferative disorders and pre-neoplastic conditions, such as, for example, hyperplasia, metaplasia, and/or	dysplasia. Highly preferred indications also include arterial disease, such as, atherosclerosis, hypertension, coronary artery disease, inflammatory vasculitides, Reynaud's disease and Reynaud's phenomenom, aneurysms, restenosis; venous and lymphatic disorders such as thrombophlebitis, lymphangitis, and lymphedema; and other vascular disorders such as peripheral vascular disease, and cancer. Highly preferred indications also	include trauma such as wounds, burns, and injured tissue (e.g., vascular injury such as, injury resulting from balloon angioplasty, and atheroschlerotic lesions), implant fixation, scarring, ischemia reperfusion injury, rheumatoid arthritis, cerebrovascular disease, renal diseases such as acute renal failure, and osteoporosis. Additional highly preferred indications include stroke, graft rejection, diabetic or other retinopathies, thrombotic and coagulative disorders, vascularitis, lymph angiogenesis, sexual disorders, age-related macular degeneration, and treatment /prevention of endometriosis and related conditions.  Additional highly preferred indications include fibromas,	heart disease, cardiac arrest, heart valve disease, and vascular disease.  Preferred indications include blood disorders (e.g., as described below under "Immune Activity", "Blood-Related Disorders", and/or "Cardiovascular Disorders"). Preferred indications include autoimmune diseases (e.g., rheumatoid arthritis, systemic lupus erythematosis, multiple sclerosis and/or as described below) and immunodeficiencies (e.g., as described below) Additional preferred indications include inflammation and inflammatory disorders (such as acute and chronic inflammatory diseases, e.g., inflammatory bowel disease
lymphangi include ca pancreatic cancer. Pre dysprolifes such as, fo	dysplasia. Highly arterial disease, such coronary artery disease are aneurysms, restenosis such as thrombophle and other vascular disease, and cancer.	include tra (e.g., vasc angioplast fixation, so arthritis, co acute rena preferred i diabetic or disorders, disorders, /preventional	heart disease, car vascular disease. disorders (e.g., as Activity", "Blooc" "Cardiovascular Jautoimmune diselupus erythemato below) and immu Additional prefer inflammatory dise

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					and Crohn's disease), and pain management.
4	HACAB68	518	Activation of Skeletal	Kinase assay. Kinase assays, for example	A highly preferred embodiment of the invention
			Mucle Cell PI3 Kinase	an GSK-3 kinase assay, for PI3 kinase	includes a method for increasing muscle cell survival An
			Signalling Pathway	signal transduction that regulate glucose	alternative highly preferred embodiment of the invention
				metabolism and cell survivial are well-	includes a method for decreasing muscle cell survival.
				known in the art and may be used or	A preferred embodiment of the invention includes a
				routinely modified to assess the ability of	method for stimulating muscle cell proliferation. In a
				polypeptides of the invention (including	specific embodiment, skeletal muscle cell proliferation is
				antibodies and agonists or antagonists of	stimulated. An alternative highly preferred embodiment of
				the invention) to promote or inhibit	the invention includes a method for inhibiting muscle cell
				glucose metabolism and cell survival.	proliferation. In a specific embodiment, skeletal muscle
				Exemplary assays for PI3 kinase activity	cell proliferation is inhibited. A preferred embodiment
				that may be used or routinely modified to	of the invention includes a method for stimulating muscle
				test PI3 kinase-induced activity of	cell differentiation. In a specific embodiment, skeletal
				polypeptides of the invention (including	muscle cell differentiation is stimulated. An alternative
				antibodies and agonists or antagonists of	highly preferred embodiment of the invention includes a
				the invention) include assays disclosed in	method for inhibiting muscle cell differentiation. In a
				Forrer et al., Biol Chem 379(8-9):1101-	specific embodiment, skeletal muscle cell differentiation is
				1110 (1998); Nikoulina et al., Diabetes	inhibited. Highly preferred indications include disorders
				49(2):263-271 (2000); and Schreyer et al.,	of the musculoskeletal system. Preferred indications
				Diabetes 48(8):1662-1666 (1999), the	include neoplastic diseases (e.g., as described below under
				contents of each of which are herein	"Hyperproliferative Disorders"), endocrine disorders (e.g.,
				incorporated by reference in its entirety.	as described below under "Endocrine Disorders"), neural
				Rat myoblast cells that may be used	disorders (e.g., as described below under "Neural Activity
				according to these assays are publicly	and Neurological Diseases"), blood disorders (e.g., as
				available (e.g., through the ATCC).	described below under "Immune Activity",
				Exemplary rat myoblast cells that may be	"Cardiovascular Disorders", and/or "Blood-Related
				used according to these assays include L6	Disorders"), immune disorders (e.g., as described below
				cells. L6 is an adherent rat myoblast cell	under "Immune Activity"), and infection (e.g., as
				line, isolated from primary cultures of rat	described below under "Infectious Disease").
	•			thigh muscle, that fuses to form	highly preferred indication is diabetes mellitus.
				multinucleated myotubes and striated	additional highly preferred indication is a complication
				fibers after culture in differentiation media.	associated with diabetes (e.g., diabetic retinopathy,
					diabetic nephropathy, kidney disease (e.g., renal failure,
					nephropathy and/or other diseases and disorders as
					described in the "Renal Disorders" section below), diabetic
					neuropaury, nerve disease and nerve damage (e.g., due to

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					diabetic neuropathy), blood vessel blockage, heart disease,
					stroke, impotence (e.g., due to diabetic neuropathy or
					blood vessel blockage), seizures, mental confusion,
					drowsiness, nonketotic hyperglycemic-hyperosmolar
					coma, cardiovascular disease (e.g., heart disease,
					atherosclerosis, microvascular disease, hypertension,
					stroke, and other diseases and disorders as described in the
					"Cardiovascular Disorders" section below), dyslipidemia,
		-			endocrine disorders (as described in the "Endocrine
					Disorders" section below), neuropathy, vision impairment
					(e.g., diabetic retinopathy and blindness), ulcers and
					impaired wound healing, infections (e.g., infectious
					diseases and disorders as described in the "Infectious
					Diseases" section below, especially of the urinary tract and
					skin), carpal tunnel syndrome and Dupuytren's
					contracture). An additional highly preferred indication
					is obesity and/or complications associated with obesity.
					Additional highly preferred indications include weight loss
					or alternatively, weight gain. Additional highly
	4				preferred indications are complications associated with
					insulin resistance. Additional highly preferred
					indications are disorders of the musculoskeletal system
					including myopathies, muscular dystrophy, and/or as
					described herein. Additional highly preferred
					indications include: myopathy, atrophy, congestive heart
					failure, cachexia, myxomas, fibromas, congenital
					o,
		**	-		heart valve disease, and vascular disease. Highly
					preferred indications include neoplasms and cancer, such
					as, rhabdomyoma, rhabdosarcoma, stomach, esophageal,
					prostate, and urinary cancer. Preferred indications also
					include breast, lung, colon, pancreatic, brain, and liver
					cancer. Other preferred indications include benign
					dysproliferative disorders and pre-neoplastic conditions,
					such as, hyperplasia, metaplasia, and/or dysplasia.
ν	HACBJ56	519	Regulation of viability	Assays for the regulation of viability and proliferation of cells in vitro are well.	A highly preferred indication is diabetes mellitus.
				איניים מונים אינים אינים אינים אינים	An additional nightly preferred illurcation is a complication

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	pancreatic beta cells.	known in the art and may be used or	associated with diabetes (e.g., diabetic retinopathy,
		routinely modified to assess the ability of	diabetic nephropathy, kidney disease (e.g., renal failure,
		polypeptides of the invention (including	nephropathy and/or other diseases and disorders as
		antibodies and agonists or antagonists of	described in the "Renal Disorders" section below), diabetic
		the invention) to regulate viability and	neuropathy, nerve disease and nerve damage (e.g., due to
		proliferation of pancreatic beta cells. For	diabetic neuropathy), blood vessel blockage, heart disease,
		example, the Cell Titer-Glo luminescent	stroke, impotence (e.g., due to diabetic neuropathy or
		cell viability assay measures the number of	blood vessel blockage), seizures, mental confusion,
		viable cells in culture based on	drowsiness, nonketotic hyperglycemic-hyperosmolar
		quantitation of the ATP present which	coma, cardiovascular disease (e.g., heart disease,
		signals the presence of metabolically	atherosclerosis, microvascular disease, hypertension,
		active cells. Exemplary assays that may be	stroke, and other diseases and disorders as described in the
		used or routinely modified to test	"Cardiovascular Disorders" section below), dyslipidemia,
		regulation of viability and proliferation of	endocrine disorders (as described in the "Endocrine
		pancreatic beta cells by polypeptides of the	Disorders" section below), neuropathy, vision impairment
		invention (including antibodies and	(e.g., diabetic retinopathy and blindness), ulcers and
		agonists or antagonists of the invention)	impaired wound healing, and infection (e.g., infectious
		include assays disclosed in: Friedrichsen	diseases and disorders as described in the "Infectious
		BN, et al., Mol Endocrinol, 15(1):136-48	Diseases" section below, especially of the urinary tract and
		(2001); Huotari MA, et al., Endocrinology,	skin), carpal tunnel syndrome and Dupuytren's
		139(4):1494-9 (1998); Hugl SR, et al., J	contracture). An additional highly preferred
		Biol Chem 1998 Jul 10;273(28):17771-9	indication is obesity and/or complications associated with
		(1998), the contents of each of which is	obesity. Additional highly preferred indications include
		herein incorporated by reference in its	weight loss or alternatively, weight gain. Aditional
		entirety. Pancreatic cells that may be used	highly preferred indications are complications associated
		according to these assays are publicly	with insulin resistance.
		available (e.g., through the ATCC) and/or	
		may be routinely generated. Exemplary	
		pancreatic cells that may be used	
		according to these assays include rat INS-1	
		cells. INS-1 cells are a semi-adherent cell	
		line established from cells isolated from an	
		X-ray induced rat transplantable	
		insulinoma. These cells retain	
		characteristics typical of native pancreatic	
		beta cells including glucose inducible	
		insulin secretion. References: Asfari et al.	

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				Endocrinology 1992 130:167	
y	HACBS22	520	Production of ICAM-1	Assays for measuring expression of	Preferred embodiments of the invention include using
)	770701111		1 127 10 110 110 110 110 110 110 110 110 110	ICAM 1 are used known in the art and	nolymentides of the invention (or antibodies agonists or
				ICAM-1 are well-known in the art and	polypeptides of the invention (of antibodies, agoinsts, of
				may be used or routinely modified to	antagonists thereot) in detection, diagnosis, prevention,
				assess the ability of polypeptides of the	and/or treatment of Vascular Disease, Atherosclerosis,
				invention (including antibodies and	Restenosis, Stroke, and Asthma.
				agonists or antagonists of the invention) to	
				regulate ICAM-1 expression. Exemplary	
				assays that may be used or routinely	
				modified to measure ICAM-1 expression	
				include assays disclosed in: Rolfe BE, et	
				al., Atherosclerosis, 149(1):99-110 (2000);	
				Panettieri RA Jr, et al., J Immunol,	
			-	154(5):2358-2365 (1995); and, Grunstein	
				MM, et al., Am J Physiol Lung Cell Mol	
				Physiol, 278(6):L1154-L1163 (2000), the	
				contents of each of which is herein	
				incorporated by reference in its entirety.	
				Cells that may be used according to these	
				assays are publicly available (e.g., through	
				the ATCC) and/or may be routinely	
				generated. Exemplary cells that may be	
				used according to these assays include	
				Aortic Smooth Muscle Cells (AOSMC):	
				such as bovine AOSMC.	
7	HADDE71	521	Activation of	Assays for the activation of transcription	A highly preferred indication is allergy. Another
			transcription through	through the Signal Transducers and	highly preferred indication is asthma. Additional
			STAT6 response	Activators of Transcription (STAT6)	highly preferred indications include inflammation and
			element in immune	response element are well-known in the art	inflammatory disorders. Preferred indications
			cells (such as natural	and may be used or routinely modified to	include blood disorders (e.g., as described below under
			killer cells).	assess the ability of polypeptides of the	"Immune Activity", "Blood-Related Disorders", and/or
				invention (including antibodies and	"Cardiovascular Disorders"). Preferred indications include
				agonists or antagonists of the invention) to	autoimmune diseases (e.g., rheumatoid arthritis, systemic
				regulate STAT6 transcription factors and	lupus erythematosis, multiple sclerosis and/or as described
				modulate the expression of multiple genes.	below) and immunodeficiencies (e.g., as described below).
				Exemplary assays for transcription through	Preferred indications include neoplastic diseases (e.g.,
				the SIAIo response element that may be	leukemia, lymphoma, melanoma, and/or as described

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				used or routinely modified to test STAT6	below under "Hyperproliferative Disorders"). Preferred
				response element activity of the	indications include neoplasms, such as, for example,
				polypeptides of the invention (including	leukemia, lymphoma, melanoma, and prostate, breast,
				antibodies and agonists or antagonists of	lung, colon, pancreatic, esophageal, stomach, brain, liver
				the invention) include assays disclosed in	and urinary cancer. Other preferred indications include
				Berger et al., Gene 66:1-10 (1998); Cullen	benign dysproliferative disorders and pre-neoplastic
				and Malm, Methods in Enzymol 216:362-	conditions, such as, for example, hyperplasia, metaplasia,
				368 (1992); Henthorn et al., Proc Natl	and/or dysplasia. Preferred indications include
				Acad Sci USA 85:6342-6346 (1988);	anemia, pancytopenia, leukopenia, thrombocytopenia,
				Georas et al., Blood 92(12):4529-4538	Hodgkin's disease, acute lymphocytic anemia (ALL),
				(1998); Moffatt et al., Transplantation	plasmacytomas, multiple myeloma, Burkitt's lymphoma,
+				69(7):1521-1523 (2000); Curiel et al., Eur	arthritis, AIDS, granulomatous disease, inflammatory
				J Immunol 27(8):1982-1987 (1997); and	bowel disease, sepsis, neutropenia, neutrophilia, psoriasis,
				Masuda et al., J Biol Chem	suppression of immune reactions to transplanted organs
				275(38):29331-29337 (2000), the contents	and tissues, hemophilia, hypercoagulation, diabetes
				of each of which are herein incorporated	mellitus, endocarditis, meningitis, and Lyme Disease.
				by reference in its entirety. T cells that	Additional preferred indications include infection (e.g., an
				may be used according to these assays are	infectious disease as described below under "Infectious
				publicly available (e.g., through the	Disease").
				ATCC). Exemplary rat natural killer cells	
				that may be used according to these assays	
				are publicly available (e.g., through the	
				ATCC).	
∞	HADDJ13	522	Activation of	Assays for the activation of transcription	ergy.
			transcription through	through the Signal Transducers and	highly preferred indication is asthma. Additional
			STAT6 response	Activators of Transcription (STAT6)	ons incl
			element in immune	response element are well-known in the art	inflammatory disorders. Preferred indications
			cells (such as natural	and may be used or routinely modified to	include blood disorders (e.g., as described below under
			killer cells).	assess the ability of polypeptides of the	"Immune Activity", "Blood-Related Disorders", and/or
				invention (including antibodies and	"Cardiovascular Disorders"). Preferred indications include
				agonists or antagonists of the invention) to	autoimmune diseases (e.g., rheumatoid arthritis, systemic
				regulate STAT6 transcription factors and	lupus erythematosis, multiple sclerosis and/or as described
				modulate the expression of multiple genes.	below) and immunodeficiencies (e.g., as described below).
				Exemplary assays for transcription through	Preferred indications include neoplastic diseases (e.g.,
				the STAT6 response element that may be	leukemia, lymphoma, melanoma, and/or as described
				used or routinely modified to test STAT6	below under "Hyperproliferative Disorders"). Preferred
				response element activity of the	indications include neoplasms, such as, for example,

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		,		polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include assays disclosed in Berger et al., Gene 66:1-10 (1998); Cullen and Malm, Methods in Enzymol 216:362-368 (1992); Henthorn et al., Proc Natl Acad Sci USA 85:6342-6346 (1988); Georas et al., Blood 92(12):4529-4538 (1998); Moffatt et al., Transplantation 69(7):1521-1523 (2000); Curiel et al., Eur J Immunol 27(8):1982-1987 (1997); and Masuda et al., J Biol Chem 275(38):29331-29337 (2000), the contents of each of which are herein incorporated by reference in its entirety. T cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary rat natural killer cells that may be used according to these assays are publicly available (e.g., through the ATCC).	leukemia, lymphoma, melanoma, and prostate, breast, lung, colon, pancreatic, esophageal, stomach, brain, liver and urinary cancer. Other preferred indications include benign dysproliferative disorders and pre-neoplastic conditions, such as, for example, hyperplasia, metaplasia, and/or dysplasia.  Preferred indications include anemia, pancytopenia, leukopenia, thrombocytopenia, Hodgkin's disease, acute lymphocytic anemia (ALL), plasmacytomas, multiple myeloma, Burkitt's lymphoma, arthritis, AIDS, granulomatous disease, inflammatory bowel disease, sepsis, neutropenia, neutrophilia, psoriasis, suppression of immune reactions to transplanted organs and tissues, hemophilia, hypercoagulation, diabetes mellitus, endocarditis, meningitis, and Lyme Disease.  Additional preferred indications include infectious Disease."
∞	HADDJ13	522	Activation of transcription through GAS response element in immune cells (such as T-cells).	Assays for the activation of transcription through the Gamma Interferon Activation Site (GAS) response element are well-known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) to regulate STAT transcription factors and modulate gene expression involved in a wide variety of cell functions. Exemplary assays for transcription through the GAS response element that may be used or routinely modified to test GAS-response element activity of polypeptides of the invention (including antibodies and agonists or	Highly preferred indications include neoplastic diseases (e.g., leukemia, lymphoma, and/or as described below under "Hyperproliferative Disorders"). Highly preferred indications include neoplasms and cancers, such as, for example, leukemia, lymphoma (e.g., T cell lymphoma, Burkitt's lymphoma, non-Hodgkins lymphoma, Hodgkin's disease), melanoma, and prostate, breast, lung, colon, pancreatic, esophageal, stomach, brain, liver and urinary cancer. Other preferred indications include benign dysproliferative disorders and pre-neoplastic conditions, such as, for example, hyperplasia, metaplasia, and/or dysplasia. Preferred indications include autoimmune diseases (e.g., rheumatoid arthritis, systemic lupus erythematosis, multiple sclerosis and/or as described below), immunodeficiencies (e.g., as described below), boosting a T cell-mediated immune response, and

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				antagonists of the invention) include assays disclosed in Berger et al., Gene 66:1-10 (1998); Cullen and Malm, Methods in Enzymol 216:362-368 (1992); Henthorn et al., Proc Natl Acad Sci USA 85:6342-6346 (1988); Matikainen et al., Blood 93(6):1980-1991 (1999); and Henttinen et al., J Immunol 155(10):4582-4587 (1995), the contents of each of which are herein incorporated by reference in its entirety. Exemplary human T cells, such as the SUPT cell line, that may be used according to these assays are publicly available (e.g., through the ATCC).	suppressing a T cell-mediated immune response.  Additional preferred indications include inflammation and inflammatory disorders. Highly preferred indications include blood disorders (e.g., as described below under "Immune Activity", "Blood-Related Disorders", and/or "Cardiovascular Disorders"), and infection (e.g., viral infections, tuberculosis, infections associated with chronic granulomatosus disease and malignant osteoporosis, and/or an infectious disease as described below under "Infectious Disease"). An additional preferred indication is idiopathic pulmonary fibrosis. Preferred indications include anemia, pancytopenia, leukopenia, thrombocytopenia, acute lymphocytic anemia (ALL), plasmacytomas, multiple myeloma, arthritis, AIDS, granulomatous disease, inflammatory bowel disease, sepsis, neutropenia, neutrophilia, psoriasis, suppression of immune reactions to transplanted organs and tissues, hemophilia, hypercoagulation, diabetes mellitus, endocarditis, meningitis, Lyme Disease, and asthma and
∞	HADDJ13	522	Activation of transcription through NFAT response element in immune cells (such as natural killer cells).	Assays for the activation of transcription through the Nuclear Factor of Activated T cells (NFAT) response element are well-known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) to regulate NFAT transcription factors and modulate expression of genes involved in immunomodulatory functions. Exemplary assays for transcription through the NFAT response element that may be used or routinely modified to test NFAT-response element activity of polypeptides of the invention (including antibodies and agonists or antagonists of the invention)	Highly preferred indications include blood disorders  (e.g., as described below under "Immune Activity", "Blood-Related Disorders", and/or "Cardiovascular Disorders"). Highly preferred indications include autoimmune diseases (e.g., rheumatoid arthritis, systemic lupus erythematosis, multiple sclerosis and/or as described below), immunodeficiencies (e.g., as described below), boosting a T cell-mediated immune response. Additional highly preferred indications include inflammation and inflammatory disorders. An additional highly preferred indication is infectious Diseases"). Preferred indications include neoplastic diseases (e.g., leukemia, lymphoma, and/or as described below under "Hyperproliferative Disorders"). Preferred indications include neoplasms and cancers, such as, for example.

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				include assays disclosed in Berger et al., Gene 66:1-10 (1998); Cullen and Malm, Methods in Enzymol 216:362-368 (1992); Henthorn et al., Proc Natl Acad Sci USA 85:6342-6346 (1988); Aramburu et al., J Exp Med 182(3):801-810 (1995); De Boer et al., Int J Biochem Cell Biol 31(10):1221-1236 (1999); Fraser et al., Eur J Immunol 29(3):838-844 (1999); and Yeseen et al., J Biol Chem 268(19):14285-14293 (1993), the contents of each of which are herein incorporated by reference in its entirety. NK cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary human NK cells that may be used according to these assays include the NK-YT cell line, which is a human natural killer cell line with cytolytic and cytotoxic	leukemia, lymphoma, and prostate, breast, lung, colon, pancreatic, esophageal, stomach, brain, liver and urinary cancer. Other preferred indications include benign dysproliferative disorders and pre-neoplastic conditions, such as, for example, hyperplasia, metaplasia, and/or dysplasia. Preferred indications also include anemia, pancytopenia, leukopenia, thrombocytopenia, Hodgkin's disease, acute lymphocytic anemia (ALL), plasmacytomas, multiple myeloma, Burkitt's lymphoma, arthritis, AIDS, granulomatous disease, inflammatory bowel disease, sepsis, neutropenia, neutrophilia, psoriasis, suppression of immune reactions to transplanted organs and tissues, hemophilia, hypercoagulation, diabetes mellitus, endocarditis, meningitis, Lyme Disease, asthma and allergy.
6	HADMB15	523	Regulation of transcription via DMEF1 response element in adipocytes and pre-adipocytes	Assays for the regulation of transcription through the DMEF1 response element are well-known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) to activate the DMEF1 response element in a reporter construct (such as that containing the GLUT4 promoter) and to regulate insulin production. The DMEF1 response element is present in the GLUT4 promoter and binds to MEF2 transcription factor and another transcription of Glut4 expression in skeletal muscle. GLUT4 is the primary insulin-responsive glucose transporter in	A highly preferred indication is diabetes mellitus. An additional highly preferred indication is a complication associated with diabetes (e.g., diabetic retinopathy, diabetic nephropathy, kidney disease (e.g., renal failure, nephropathy and/or other diseases and disorders as described in the "Renal Disorders" section below), diabetic neuropathy, nerve disease and nerve damage (e.g., due to diabetic neuropathy), blood vessel blockage, heart disease, stroke, impotence (e.g., due to diabetic neuropathy or blood vessel blockage), seizures, mental confusion, drowsiness, nonketotic hyperglycemic-hyperosmolar coma, cardiovascular disease (e.g., heart disease, atherosclerosis, microvascular disease, hypertension, stroke, and other diseases and disorders as described in the "Cardiovascular Disorders" section below), dyslipidemia, endocrine disorders (as described in the "Endocrine Disorders" section below), neuropathy, vision impairment

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				fat and muscle tissue. Exemplary assays that may be used or routinely modified to test for DMEF1 response element activity (in adipocytes and pre-adipocytes) by polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include assays disclosed in Thai, M.V., et al., J Biol Chem, 273(23):14285-92 (1998); Mora, S., et al., J Biol Chem, 269(45):28514-21 (1994); "Identification of a 30-base pair regulatory element and novel DNA binding protein that regulates the human GLUT4 promoter in transgenic mice", J Biol Chem. 2000 Aug 4;275(31):23666-73; Berger, et al., Gene 66:1-10 (1988); and, Cullen, B., et al., Methods in Enzymol. 216:362–368 (1992), the contents of each of which is herein incorporated by reference in its entirety. Adipocytes and pre-adipocytes that may be used according to these assays are publicly available (e.g., through the ATCC) and/or may be routinely generated. Exemplary	impaired wound healing, and infection (e.g., infectious diseases and disorders as described in the "Infectious Diseases" section below, especially of the urinary tract and skin), carpal tunnel syndrome and Dupuytren's Contracture).  An additional highly preferred indication is obesity and/or complications associated with obesity. Additional highly preferred indications include weight loss or alternatively, weight gain.  Additional highly preferred indications associated with insulin resistance.
				cells that may be used according to these assays include the mouse 3T3-L1 cell line which is an adherent mouse preadipocyte cell line. Mouse 3T3-L1 cells are a continuous substrain of 3T3 fibroblasts developed through clonal isolation. These cells undergo a pre-adipocyte to adiposelike conversion under appropriate differentiation culture conditions.	
6	HADMB15	523	Regulation of apoptosis of immune cells (such as mast cells).	Caspase Apoptosis. Assays for caspase apoptosis are well known in the art and may be used or routinely modified to	Preferred embodiments of the invention include using polypeptides of the invention (or antibodies, agonists, or antagonists thereof) in detection, diagnosis, prevention,

and/or treatment of asthma, allergy, hypersensitivity and inflammation.	A highly preferred embodiment of the invention includes a method for stimulating natural killer cell
assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) to regulate caspase protease-mediated apoptosis in immune cells (such as, for example, in mast cells). Mast cells are found in connective and mucosal tissues throughout the body, and their activation via immunoglobulin E-antigen, promoted by T helper cell type 2 cytokines, is an important component of allergic disease. Dysregulation of mast cell apoptosis may play a role in allergic disease and mast cell tumor survival. Exemplary assays for caspase apoptosis that may be used or routinely modified to test capase apoptosis activity induced by polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include the assays disclosed in: Masuda A, et al., J Biol Chem, 276(28):26107-26113 (2001); Yeatman CF 2nd, et al., J Exp Med, 192(8):1093-1103 (2000); Lee et al., FEBS Lett 485(2-3): 122-126 (2000); and Karsan and Harlan, J Atheroscler Thromb 3(2): 75-80 (1996); the contents of each of which are herein incorporated by reference in its entirety. Immune cells that may be used according to these assays include mast cells such as the HMC burner and the state of the such according to these assays include mast cells such as the HMC burner and the such according to these	Kinase assays, for example an Elk-1 kinase assays, for ERK signal
	Activation of Natural Killer Cell ERK
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Y. Z. Z. Z. Z. Z. Z. Z. Z. Z. Z. Z. Z. Z.	S.	transduction that regulate cell proliferation or differentiation are well known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) to	activation, and differentiation. Exemplary assays for ERK kinase activity that may be kinase-induced activity of polypeptides of agonists or antagonists of the assays disclosed in Forrer et activity of properties or antagonists of the invention (include the assays disclosed in Forrer et activity).	al., Biol Chem 379(8-9):1101-1110 (1998); Kyriakis JM, Biochem Soc Symp (4:29-48 (1999); Chang and Karin, Nature (4:29-48 (1999); Chang and Karin, Nature (4:29-48 (1999); Chang and Karin, Nature (4:29-48 (1999); Chang and Karin, Nature (4:29-48 (1999); Chang and Karin, Nature (4:29-48 (1999); Chang and Karin, Nature (4:29-48 (1999); Chang and Karin, Nature (4:29-48 (1999); Chang and Karin, Nature (4:29-48 (1999); Chang and Karin, Nature (5:29-48 (1999); Chang and Karin, Nature (6:29-	>
			PT US BS DE PT VI PT VI PT PT PT PT PT PT PT PT PT PT PT PT PT	20 (1) P. P. P.	3 E & & X

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A highly preferred embodiment of the invention includes a method for stimulating the production of IFNg. An alternative highly preferred embodiment of the invention includes a method for inhibiting the production of IFNg. Highly preferred indications include blood disorders (e.g., as described below under "Immune Activity" "Blood-Related Disorders" and/or	"Cardiovascular Disorders"), and infection (e.g., viral infections, tuberculosis, infections associated with chronic granulomatosus disease and malignant osteoporosis, and/or as described below under "Infectious Disease"). Highly preferred indications include autoimmune disease (e.g., rheumatoid arthritis, systemic lupus erythematosis, multiple sclerosis and/or as described below), hoosting a Trimmunodeficiency (e.g., as described below) hoosting a Trimmunodeficiency (e.g., as described below).	minimulocation (e.g., as used the probability), boosting a reell-mediated immune response, and suppressing a T cell-mediated immune response. Additional highly preferred indications include inflammation and inflammatory disorders. Additional preferred indications include idiopathic pulmonary fibrosis. Highly preferred indications include neoplastic diseases (e.g., leukemia, lymphoma, melanoma, and/or as described below under "Hyperproliferative Disorders"). Highly preferred	indications include neoplasms and cancers, such as, for example, leukemia, lymphoma, melanoma, and prostate, breast, lung, colon, pancreatic, esophageal, stomach, brain, liver and urinary cancer. Other preferred indications include benign dysproliferative disorders and preneoplastic conditions, such as, for example, hyperplasia, metaplasia, and/or dysplasia. Preferred indications include anemia, pancytopenia, leukopenia,	thrombocytopenia, Hodgkin's disease, acute lymphocytic anemia (ALL), plasmacytomas, multiple myeloma, Burkitt's lymphoma, arthritis, AIDS, granulomatous disease, inflammatory bowel disease, sepsis, neutropenia, neutrophilia, psoriasis, suppression of immune reactions to transplanted organs and tissues, hemophilia,
IFNgamma FMAT. IFNg plays a central role in the immune system and is considered to be a proinflammatory cytokine. IFNg promotes TH1 and inhibits TH2 differentiation; promotes IgG2a and inhibits IgE secretion; induces marronhage activation; and increases	MHC expression. Assays for immunomodulatory proteins produced by T cells and NK cells that regulate a variety of inflammatory activities and inhibit TH2 helper cell functions are well known in the art and may be used or routinely modified to assess the ability of polypeptides of the invastion (including anti-hodies and	agonists or antagonists of the invention) to mediate immunomodulation, regulate inflammatory activities, modulate TH2 helper cell function, and/or mediate humoral or cell-mediated immunity. Exemplary assays that test for immunomodulatory proteins evaluate the production of cytokines, such as Interferon	gamma (IFNg), and the activation of T cells. Such assays that may be used or routinely modified to test immunomodulatory activity of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include the assays disclosed in Miraglia et al., J Biomolecular	Screening 4:193-204 (1999); Rowland et al., "Lymphocytes: a practical approach" Chapter 6:138-160 (2000); Gonzalez et al., J Clin Lab Anal 8(5):225-233 (1995); Billiau et al., Ann NY Acad Sci 856:22-32 (1998); Boehm et al., Annu Rev Immunol
Production of IFNgamma using a T cells				
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			15:749-795 (1997), and Rheumatology (Oxford) 38(3):214-20 (1999), the contents of each of which are herein incorporated	hypercoagulation, diabetes mellitus, endocarditis, meningitis, Lyme Disease, asthma and allergy.
			by reference in its entirety. Human T cells that may be used according to these assays	
			may be isolated using techniques disclosed herein or otherwise known in the art.	
			Human T cells are primary human	
			lymphocytes that mature in the thymus and	
			or CD8. These cells mediate humoral or	
			cell-mediated immunity and may be	
			preactivated to enhance responsiveness to	
OCHICATI	202	TT and lotion of LIT A	Immunomodulatory lactors.	Highly preferred indications include blood disorders
II HAGDW20	272	DP and activation of T	for correct presentation of antigen to CD4+	(e.g., as described below under "Immune Activity",
		cells	T cells. Deregulation of MHC class II has	"Blood-Related Disorders", and/or "Cardiovascular
			been associated with autoimmune diseases	Disorders"). Highly preferred indications include
			(e.g., diabetes, rheumatoid arthritis,	autoimmune diseases (e.g., rheumatoid arthritis, systemic
			systemic lupus erythematosis, and multiple	lupus erythematosis, multiple sclerosis and/or as described
			sclerosis). Assays for immunomodulatory	below) and immunodeficiencies (e.g., as described below),
			proteins expressed on MHC class II	boosting a T cell-mediated immune response, and
			expressing T cells and antigen presenting	ely, su
			cells are well known in the art and may be	
			used or routinely modified to assess the	mellitus. An additional highly preferred indication
			ability of polypeptides of the invention	is a complication associated with diabetes (e.g., diabetic
			(including antibodies and agonists or	retinopathy, diabetic nephropathy, kidney disease (e.g.,
			antagonists of the invention) to modulate	renal failure, nephropathy and/or other diseases and
			the activation of 1 cells, and/or mediate	disorders as described in the Including Disorders section
			humoral or cell-mediated immunity.	demonstrate discussional versions and inclive described the second demonstrate of the description of the des
			Exemplary assays that test lor	dalliage (e.g., due to diabetic liculopanily), blood vessel
			immunomodulatory proteins evaluate the	blockage, heart disease, stroke, impotence (e.g., due to
			upregulation of MHC class II products,	diabetic neuropathy or blood vessel blockage), seizures,
			such as HLA-DR antigens, and the	mental confusion, drowsiness, nonketotic hyperglycemic-
			activation of T cells. Such assays that may	hyperosmolar coma, cardiovascular disease (e.g., heart
			be used or routinely modified to test	disease, atherosclerosis, microvascular disease,
			immunomodulatory activity of	hypertension, stroke, and ounce diseases and disorders as

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12	HAGEG10	526	Activation of	Assays for the activation of transcription	meningitis, Lyme Disease, inflammation and inflammatory disorders, and asthma and allergy.  Highly preferred indications include neoplastic diseases feellelsemia, lymphoma, and/or as described below
			GAS response element in immune cells (such	Site (GAS) response element are well- known in the art and may be used or	under "Hyperproliferative Disorders"). Highly preferred indications include neoplasms and cancers, such as, for
				routinely modified to assess the ability of	example, leukemia, lymphoma (e.g., T cell lymphoma, Burkitt's lymphoma, non-Hodgkins lymphoma, Hodgkin's
				antibodies and agonists or antagonists of	disease), melanoma, and prostate, breast, lung, colon,
				the invention) to regulate STAT transcription factors and modulate gene	pancreatic, esophageal, stomach, brain, liver and urinary cancer. Other preferred indications include benign
				expression involved in a wide variety of	dysproliferative disorders and pre-neoplastic conditions,
				cell functions. Exemplary assays for	such as, for example, hyperplasia, metaplasia, and/or dyenlasia — Preferred indications include autoimmune
				element that may be used or routinely	diseases (e.g., rheumatoid arthritis, systemic lupus
				modified to test GAS-response element	erythematosis, multiple sclerosis and/or as described
				activity of polypeptides of the invention	below), immunodeficiencies (e.g., as described below),
				(including antibodies and agonists or	boosting a T cell-mediated immune response, and
				antagonists of the invention) include	suppressing a 1 cell-mediated immune response. Additional preferred indications include inflammation and
				assays disclosed in Derger et al., Celle 66:1-10 (1998): Cullen and Malm.	inflammatory disorders. Highly preferred indications
				Methods in Enzymol 216:362-368 (1992);	include blood disorders (e.g., as described below under
				Henthorn et al., Proc Natl Acad Sci USA	"Immune Activity", "Blood-Related Disorders", and/or
				85:6342-6346 (1988); Matikainen et al.,	"Cardiovascular Disorders"), and infection (e.g., viral
				Blood 93(6):1980-1991 (1999); and	infections, tuberculosis, infections associated with chronic
			-	Henttinen et al., J Immunol 155(10):4582-	granulomatosus disease and malignant osteoporosis,
				4587 (1995), the contents of each of which	and/or an infectious disease as described below under
				are nerein incorporated by reference in its	is idionathic nilmonary fibrosis.  Preferred indications
				may be used according to these assays are	ıkol
				publicly available (e.g., through the	thrombocytopenia, acute lymphocytic anemia (ALL),
				ATCC). Exemplary T cells that may be	plasmacytomas, multiple myeloma, arthritis, AIDS,
				used according to these assays include the	granulomatous disease, inflammatory bowel disease,
				CTLL cell line, which is a suspension	sepsis, neutropenia, neutrophilia, psoriasis, suppression of
				culture of IL-2 dependent cytotoxic 1	Immune reactions to transplanted organs and ussues,
				cells.	nemophilia, hypercoagulation, diabetes metilitus, endocarditis meninoitis I vme Disease, and asthma and

					allarmy
					alicigy.
13	HAGE079	527	Activation of	Assays for the activation of transcription	Highly preferred indications include neoplastic diseases
	,		transcription through	through the Gamma Interferon Activation	(e.g., leukemia, lymphoma, and/or as described below
	٠		GAS response element	Site (GAS) response element are well-	under "Hyperproliferative Disorders"). Highly preferred
			in immine cells (such	known in the art and may be used or	indications include neoplasms and cancers, such as, for
			as T-cells)	routinely modified to assess the ability of	example, leukemia, lymphoma (e.g., T cell lymphoma,
			./6	polyneptides of the invention (including	Burkitt's lymphoma, non-Hodgkins lymphoma, Hodgkin's
				antibodies and agonists or antagonists of	disease), melanoma, and prostate, breast, lung, colon,
				the invention) to regulate STAT	pancreatic, esophageal, stomach, brain, liver and urinary
				transcription factors and modulate gene	cancer. Other preferred indications include benign
				expression involved in a wide variety of	dysproliferative disorders and pre-neoplastic conditions,
				cell functions. Exemplary assays for	such as, for example, hyperplasia, metaplasia, and/or
				transcription through the GAS response	dysplasia. Preferred indications include autoimmune
				element that may be used or routinely	diseases (e.g., rheumatoid arthritis, systemic lupus
				modified to test GAS-response element	erythematosis, multiple sclerosis and/or as described
				activity of polypeptides of the invention	below), immunodeficiencies (e.g., as described below),
				(including antibodies and agonists or	boosting a T cell-mediated immune response, and
				antagonists of the invention) include	suppressing a T cell-mediated immune response.
				assays disclosed in Berger et al., Gene	Additional preferred indications include inflammation and
				66:1-10 (1998); Cullen and Malm,	inflammatory disorders. Highly preferred indications
				Methods in Enzymol 216:362-368 (1992);	include blood disorders (e.g., as described below under
				Henthorn et al., Proc Natl Acad Sci USA	"Immune Activity", "Blood-Related Disorders", and/or
				85:6342-6346 (1988); Matikainen et al.,	"Cardiovascular Disorders"), and infection (e.g., viral
				Blood 93(6):1980-1991 (1999); and	infections, tuberculosis, infections associated with chronic
				Henttinen et al., J Immunol 155(10):4582-	granulomatosus disease and malignant osteoporosis,
				4587 (1995), the contents of each of which	and/or an infections disease as described below under
				are herein incorporated by reference in its	onal
				entirety. Exemplary mouse T cells that	is idiopathic pulmonary fibrosis. Preferred indications
				may be used according to these assays are	include anemia, pancytopenia, leukopenia,
				publicly available (e.g., through the	thrombocytopenia, acute lymphocytic anemia (ALL),
				ATCC). Exemplary T cells that may be	plasmacytomas, multiple myeloma, arthritis, AIDS,
				used according to these assays include the	granulomatous disease, inflammatory bowel disease,
				CTLL cell line, which is a suspension	sepsis, neutropenia, neutrophilia, psoriasis, suppression of
				culture of IL-2 dependent cytotoxic T	immune reactions to transplanted organs and tissues,
				cells.	hemophilia, hypercoagulation, diabetes mellitus,
					endocarditis, meningitis, Lyme Disease, and asthma and
					anergy.

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Highly preferred indications include blood disorders (e.g., as described below under "Immune Activity", "Blood-Related Disorders", and/or "Cardiovascular Disorders"). Highly preferred indications include autoimmune diseases (e.g., rheumatoid arthritis, systemic lupus erythematosis, multiple sclerosis and/or as described below) and immunodeficiencies (e.g., as described below),	alternatively, suppressing a T cell-mediated immune response.  A highly preferred indication is diabetes mellitus.  An additional highly preferred indication is a complication associated with diabetes (e.g., diabetic retinopathy, diabetic nephropathy, kidney disease (e.g., renal failure, nephropathy and/or other diseases and disorders as described in the "Renal Disorders" section below), diabetic neuropathy, nerve disease and nerve damage (e.g., due to diabetic neuropathy), blood vessel blockage, heart disease, stroke, impotence (e.g., due to diabetic neuropathy or blood vessel blockage), seizures, mental confusion, drowsiness, nonketotic hyperglycemichyperosmolar coma, cardiovascular disease (e.g., heart disease, atherosclerosis, microvascular disease, heart disease, and other diseases and disorders as described in the "Cardiovascular Disorders" section below), dyslipidemia, endocrine disorders (as described in	the "Endocrine Disorders" section below), neuropathy, vision impairment (e.g., diabetic retinopathy and blindness), ulcers and impaired wound healing, and infection (e.g., infectious diseases and disorders as described in the "Infectious Diseases" section below, especially of the urinary tract and skin), carpal tunnel syndrome and Dupuytren's contracture). An additional highly preferred indication is obesity and/or complications associated with obesity. Additional highly preferred indications include weight loss or alternatively, weight gain. Additional highly preferred indications associated with insulin resistance.
sential CD4+ II has seases wiltiple atory	expressing T cells and antigen presenting cells are well known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) to modulate the activation of T cells, and/or mediate humoral or cell-mediated immunity.  Exemplary assays that test for immunomodulatory proteins evaluate the upregulation of MHC class II products, such as HLA-DR antigens, and the activation of T cells. Such assays that may be used or routinely modified to test immunomodulatory activity of polypeptides of the invention (including antibodies and agonists or antagonists of	la, la,
Upregulation of HLA-DR and activation of T cells		
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## DOOMDOOM . DOLLOOK

HA	HAGFS57 528	Proliferation, differentiation, and/or for cytokine production in immune cells (such as king immune cells).  Proliferation, and/or for cytokine production in (in immune cells).  Proliferation, and/or for cytokine production in (in immune cells).	Human T cells that may be used according to these assays may be isolated using techniques disclosed herein or otherwise known in the art. Human T cells are primary human lymphocytes that mature in the thymus and express a T Cell receptor and CD3, CD4, or CD8. These cells mediate humoral or cell-mediated immunity and may be preactivated to enhance responsiveness to immunomodulatory factors.  Kinase assays, for example kinase assays for members of the MAP kinase family (including p38, JAK, and ERK) are well known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) to promote or inhibit cell proliferation, differentiation, and/or cytokine production in immune cells such as T-cells. Exemplary assays for MAP	Additonal highly preferred indications are disorders of the musculoskeletal systems including myopathies, muscular dystrophy, and/or as described herein.  An additional preferred indication is infection (e.g., AIDS, and/or as described below under "Infectious Disease").  Preferred indications include endocrine disorders (e.g., as described below under "Endocrine Disorders"). and neoplastic diseases (e.g., leukemia, lymphoma, and/or as described below under "Hyperproliferative Disorders").  Preferred indications include neoplasms and cancer, such as, for example, leukemia, lymphoma, and prostate, breast, lung, colon, pancreatic, esophageal, stomach, brain, liver and urinary cancer. Other preferred indications include benign dysproliferative disorders and pre-neoplastic conditions, such as, for example, hyperplasia, metaplasia, and/or dysplasia.  Preferred indications also include anemia, pancytopenia, leukopenia, thrombocytopenia, leukopenia, thrombocytopenia, leukopenia, thrombocytopenia, hypercoagulation, endocarditis, blasmacytomas, multiple myeloma, Burkitt's lymphoma, and tissues, hemophilia, hypercoagulation, endocarditis, meningitis, Lyme Disease, inflammation and inflammatory disorders, and asthma and allergy.  Preferred embodiments of the invention include using polypeptides of the invention (or antibodies, agonists, or antagonists thereof) in detection, diagnosis, prevention, and/or treatment of inflammation, infection, allergy, asthma, autoimmunity, and cancer.
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e e W. des	T cells  A highly preferred embodiment of the invention includes a method for stimulating (e.g., increasing) IL-6 production. An alternative highly preferred embodiment of the invention includes a method for inhibiting (e.g., and the invention includes a method for inhibiting (e.g., and indication is the stimulation or enhancement of mucosal immunity. Highly preferred indications include blood disorders (e.g., as described below under "Infectious Disease"). Highly preferred indications include autoimmune diseases (e.g., rheumatoid arthritis, systemic lupus erythematosis, multiple sclerosis and/or as described below) and immunodeficiencies (e.g., as described below) and immunodeficiencies (e.g., as described below). Highly preferred indications also include boosting a B cellmediated immune response. Highly preferred indications include inflammation and inflammatory disorders. Additional highly preferred indications include asthma and allergy. Highly preferred indications include neoplastic diseases (e.g., myeloma, plasmacytoma, leukemia, lymphoma, melanoma, and/or as described
or routinely modified to test polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include the assays disclosed in: Rincon M., Curr Opin Immunol; 13(3):339-345 (2001); Wang LH, et al., J Immunol, 162(7):3897-3904 (1999); Sakamoto H, et al., J Biol Chem, 275(46):35857-35862 (2000), the contents of each of which are herein incorporated by reference in its entirety. Exemplary immune cells (for example, T-cells) that may be used according to these assays include the mouse CTLL cell line.	IL-6 FMAT. IL-6 is produced by T cells and has strong effects on B cells. IL-6 participates in IL-4 induced IgE production and increases IgA production (IgA plays a role in mucosal immunity). IL-6 induces cytotoxic T cells. Deregulated expression of IL-6 has been linked to autoimmune disease, plasmacytomas, myelomas, and chronic hyperproliferative diseases.  Assays for immunomodulatory and differentiation factor proteins produced by a large variety of cells where the expression level is strongly regulated by cytokines, growth factors, and hormones are well known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention and function modulate T cell proliferation and function. Exemplary assays that test for immunomodulatory proteins evaluate the
	Production of IL-6
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				production of cytokines, such as IL-6, and	below under "Hyperproliferative Disorders"). Highly
				the stimulation and upregulation of T cell	preferred indications include neoplasms and cancers, such
				proliferation and functional activities.	as, myeloma, plasmacytoma, leukemia, lymphoma,
				Such assays that may be used or routinely	melanoma, and prostate, breast, lung, colon, pancreatic,
				modified to test immunomodulatory and	esophageal, stomach, brain, liver and urinary cancer.
				diffferentiation activity of polypeptides of	Other preferred indications include benign dysproliferative
				the invention (including antibodies and	disorders and pre-neoplastic conditions, such as, for
				agonists or antagonists of the invention)	example, hyperplasia, metaplasia, and/or dysplasia.
				include assays disclosed in Miraglia et al.,	Preferred indications include anemia, pancytopenia,
				J Biomolecular Screening 4:193-	leukopenia, thrombocytopenia, Hodgkin's disease, acute
				204(1999); Rowland et al., "Lymphocytes:	lymphocytic anemia (ALL), multiple myeloma, Burkitt's
				a practical approach" Chapter 6:138-160	lymphoma, arthritis, AIDS, granulomatous disease,
		_		(2000); and Verhasselt et al., J Immunol	inflammatory bowel disease, sepsis, neutropenia,
				158:2919-2925 (1997), the contents of	neutrophilia, psoriasis, suppression of immune reactions to
				each of which are herein incorporated by	transplanted organs and tissues, hemophilia,
				reference in its entirety. Human dendritic	hypercoagulation, diabetes mellitus, endocarditis,
				cells that may be used according to these	meningitis, and Lyme Disease. An additonal preferred
				assays may be isolated using techniques	indication is infection (e.g., an infectious disease as
				disclosed herein or otherwise known in the	described below under "Infectious Disease").
				art. Human dendritic cells are antigen	
				presenting cells in suspension culture,	
				which, when activated by antigen and/or	
				cytokines, initiate and upregulate T cell	
				proliferation and functional activities.	
15	HAGHN57	529	Activation of	Assays for the activation of transcription	A preferred embodiment of the invention includes a
			transcription through	through the Serum Response Element	method for inhibiting (e.g., reducing) TNF alpha
			serum response element	(SRE) are well-known in the art and may	production. An alternative highly preferred embodiment of
			in immune cells (such	be used or routinely modified to assess the	the invention includes a method for stimulating (e.g.,
			as T-cells).	ability of polypeptides of the invention	increasing) TNF alpha production. Preferred indications
				(including antibodies and agonists or	include blood disorders (e.g., as described below under
				antagonists of the invention) to bind the	"Immune Activity", "Blood-Related Disorders", and/or
				serum response factor and modulate the	"Cardiovascular Disorders"), Highly preferred indications
				expression of genes involved in growth	include autoimmune diseases (e.g., rheumatoid arthritis,
				and upregulate the function of growth-	systemic lupus erythematosis, Crohn's disease, multiple
				related genes in many cell types.	sclerosis and/or as described below), immunodeficiencies
				Exemplary assays for transcription through	(e.g., as described below), boosting a T cell-mediated
				the SKE that may be used or routinely	immune response, and suppressing a 1 cell-mediated

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				modified to test SRE activity of the	immune response. Additional highly preferred indications
				polypeptides of the invention (including antibodies and agonists or antagonists of	include initalination and initalinately disorders, and treating joint damage in patients with rheumatoid arthritis.
				the invention) include assays disclosed in	An additional highly preferred indication is sepsis.
				Berger et al., Gene 66:1-10 (1998); Cullen	Highly preferred indications include neoplastic diseases
				and Malm, Methods in Enzymol 216:362-	(e.g., leukemia, lymphoma, and/or as described below
				368 (1992); Henthorn et al., Proc Natl	under "Hyperproliferative Disorders"). Additionally,
				Acad Sci USA 85:6342-6346 (1988);	highly preferred indications include neoplasms and
				Benson et al., J Immunol 153(9):3862-	cancers, such as, for example, leukemia, lymphoma,
				3873 (1994); and Black et al., Virus Genes	melanoma, glioma (e.g., malignant glioma), solid tumors,
				12(2):105-117 (1997), the content of each	and prostate, breast, lung, colon, pancreatic, esophageal,
			,	of which are herein incorporated by	stomach, brain, liver and urinary cancer. Other preferred
				reference in its entirety. T cells that may	indications include benign dysproliferative disorders and
				be used according to these assays are	=
				publicly available (e.g., through the	hyperplasia, metaplasia, and/or dysplasia. Preferred
				ATCC). Exemplary human T cells, such	indications include anemia, pancytopenia, leukopenia,
				as the MOLT4, that may be used according	thrombocytopenia, Hodgkin's disease, acute lymphocytic
				to these assays are publicly available (e.g.,	anemia (ALL), plasmacytomas, multiple myeloma,
				through the ATCC).	Burkitt's lymphoma, arthritis, AIDS, granulomatous
					disease, inflammatory bowel disease, neutropenia,
					neutrophilia, psoriasis, suppression of immune reactions to
					transplanted organs and tissues, hemophilia,
					hypercoagulation, diabetes mellitus, endocarditis,
					meningitis, Lyme Disease, cardiac reperfusion injury, and
					asthma and allergy. An additional preferred indication
					is infection (e.g., an infectious disease as described below
31	UACUNS7	500	Activition	Account for the cotion of two accomination	A bight macformed indication is allower.
3	CNIEDAII	676	Activation of	Assays for the activation of transcription	cigy.
			uanscripuon unougn	unough une Signar Transducers and	Ingiliy preferred indication is asumia. Additional
			STAT6 response	Activators of Transcription (STAT6)	ons incl
			element in immune	response element are well-known in the art	inflammatory disorders. Preferred indications
			cells (such as natural	and may be used or routinely modified to	include blood disorders (e.g., as described below under
			killer cells).	assess the ability of polypeptides of the	"Immune Activity", "Blood-Related Disorders", and/or
				invention (including antibodies and	"Cardiovascular Disorders"). Preferred indications include
				agonists or antagonists of the invention) to	autoimmune diseases (e.g., rheumatoid arthritis, systemic
				regulate STAT6 transcription factors and	lupus erythematosis, multiple sclerosis and/or as described
				modulate the expression of multiple genes.	below) and immunodeficiencies (e.g., as described below).

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				Exemplary assays for transcription through	Preferred indications include neoplastic diseases (e.g.,
				the STAT6 response element that may be	leukemia, lymphoma, melanoma, and/or as described
				used or routinely modified to test STAT6	below under "Hyperproliferative Disorders"). Preferred
				response element activity of the	indications include neoplasms, such as, for example,
				polypeptides of the invention (including	leukemia, lymphoma, melanoma, and prostate, breast,
				antibodies and agonists or antagonists of	lung, colon, pancreatic, esophageal, stomach, brain, liver
				the invention) include assays disclosed in	and urinary cancer. Other preferred indications include
				Berger et al., Gene 66:1-10 (1998); Cullen	benign dysproliferative disorders and pre-neoplastic
				and Malm, Methods in Enzymol 216:362-	as, for examp
				368 (1992); Henthorn et al., Proc Natl	and/or dysplasia. Preferred indications include
				Acad Sci USA 85:6342-6346 (1988);	anemia, pancytopenia, leukopenia, thrombocytopenia,
				Georas et al., Blood 92(12):4529-4538	Hodgkin's disease, acute lymphocytic anemia (ALL),
				(1998); Moffatt et al., Transplantation	plasmacytomas, multiple myeloma, Burkitt's lymphoma,
				69(7):1521-1523 (2000); Curiel et al., Eur	arthritis, AIDS, granulomatous disease, inflammatory
				J Immunol 27(8):1982-1987 (1997); and	bowel disease, sepsis, neutropenia, neutrophilia, psoriasis,
				Masuda et al., J Biol Chem	suppression of immune reactions to transplanted organs
				275(38):29331-29337 (2000), the contents	and tissues, hemophilia, hypercoagulation, diabetes
				of each of which are herein incorporated	mellitus, endocarditis, meningitis, and Lyme Disease.
				by reference in its entirety. T cells that	Additional preferred indications include infection (e.g., an
				may be used according to these assays are	infectious disease as described below under "Infectious
				publicly available (e.g., through the	Disease").
				ATCC). Exemplary rat natural killer cells	
				that may be used according to these assays	
				are publicly available (e.g., through the	
				ATCC).	
15	HAGHIN57	529	Activation of	Assays for the activation of transcription	Highly preferred indications include blood disorders
			transcription through	through the Nuclear Factor of Activated T	(e.g., as described below under "Immune Activity",
			NFAT response	cells (NFAT) response element are well-	"Blood-Related Disorders", and/or "Cardiovascular
			element in immune	known in the art and may be used or	Disorders"). Highly preferred indications include
			cells (such as natural	routinely modified to assess the ability of	autoimmune diseases (e.g., rheumatoid arthritis, systemic
			killer cells).	polypeptides of the invention (including	lupus erythematosis, multiple sclerosis and/or as described
				antibodies and agonists or antagonists of	below), immunodeficiencies (e.g., as described below),
				the invention) to regulate NFAT	boosting a T cell-mediated immune response, and
				transcription factors and modulate	suppressing a T cell-mediated immune response.
				expression of genes involved in	Additional highly preferred indications include
				immunomodulatory functions. Exemplary	inflammation and inflammatory disorders. An additional
				assays for transcription through the INFA1	nignly preferred indication is infection (e.g., an infectious

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			the invention) to modulate the activation of	and/or inactivation NK cells Highly preferred
			T cells, and/or mediate humoral or cell-	tion
			mediated immunity. Exemplary assays	e. 69.
			that test for immunomodulatory proteins	Activity"). Preferred indications include blood
			evaluate the upregulation of cell surface	disorders (e.g., as described below under "Immune
			markers, such as CD69, and the activation	Activity", "Blood-Related Disorders", and/or
			of T cells. Such assays that may be used	"Cardiovascular Disorders"). Highly preferred indications
			or routinely modified to test	include autoimmune diseases (e.g., rheumatoid arthritis,
			immunomodulatory activity of	systemic lupus erythematosis, multiple sclerosis and/or as
			polypeptides of the invention (including	described below), immunodeficiencies (e.g., as described
	-		antibodies and agonists or antagonists of	below), boosting a T cell-mediated immune response and
			the invention) include, for example, the	alternatively suppressing a T cell-mediated immune
			assays disclosed in Miraglia et al., J	response, and boosting a B cell-mediated immune
			Biomolecular Screening 4:193-204 (1999);	response and alternatively suppressing a B cell-mediated
			Rowland et al., "Lymphocytes: a practical	immune response. An additional highly preferred
			approach" Chapter 6:138-160 (2000);	indication includes infection (e.g., as described below
			Ferenczi et al., J Autoimmun 14(1):63-78	under "Infectious Disease"). Preferred indications also
			(200); Werfel et al., Allergy 52(4):465-469	include anemia, pancytopenia, leukopenia,
			(1997); Taylor-Fishwick and Siegel, Eur J	thrombocytopenia, Hodgkin's disease, acute lymphocytic
			Immunol 25(12):3215-3221 (1995); and	anemia (ALL), plasmacytomas, multiple myeloma,
			Afetra et al., Ann Rheum Dis 52(6):457-	Burkitt's lymphoma, arthritis, AIDS, granulomatous
			460 (1993), the contents of each of which	disease, inflammatory bowel disease, sepsis, neutropenia,
			are herein incorporated by reference in its	neutrophilia, psoriasis, suppression of immune reactions to
			entirety. Human T cells that may be used	transplanted organs and tissues, hemophilia,
			according to these assays may be isolated	hypercoagulation, diabetes mellitus, endocarditis,
			using techniques disclosed herein or	meningitis, Lyme Disease, inflammation and
			otherwise known in the art. Human T cells	inflammatory disorders, asthma, and allergies.
			are primary human lymphocytes that	Preferred indications also include neoplastic diseases (e.g.,
			mature in the thymus and express a T Cell	leukemia, lymphoma, and/or as described below under
•			receptor and CD3, CD4, or CD8. These	"Hyperproliferative Disorders"). Preferred indications
			cells mediate humoral or cell-mediated	include neoplasms, such as, for example, leukemia,
			immunity and may be preactivated to	lymphoma, and prostate, breast, lung, colon, pancreatic,
			enhance responsiveness to	esophageal, stomach, brain, liver and urinary cancer.
			immunomodulatory factors.	Other preferred indications include benign dysproliferative
				disorders and pre-neoplastic conditions, such as, for
17 HAJAA47	531	Production of TNF	TNFa FMAT. Assays for	A highly preferred embodiment of the invention

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		inflammation and inflammatory disorders, and treating joint damage in patients with rheumatoid arthritis. An additional highly preferred indication is sepsis. Highly preferred indications include neoplastic diseases (e.g., leukemia, lymphoma, and/or as described below under "Hyperproliferative Disorders"). Additionally, highly preferred indications include neoplasms and cancers, such as, leukemia, lymphoma, melanoma, glioma (e.g., malignant glioma), solid tumors, and prostate, breast,	
immunomodulatory proteins produced by activated macrophages, T cells, fibroblasts, smooth muscle, and other cell types that exert a wide variety of inflammatory and cytotoxic effects on a variety of cells are well known in the art and may be used or	routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) to mediate immunomodulation, modulate inflammation and cytotoxicity. Exemplary assays that test for immunomodulatory proteins evaluate the production of	cytokines such as tumor necrosis factor alpha (TNFa), and the induction or inhibition of an inflammatory or cytotoxic response. Such assays that may be used or routinely modified to test immunomodulatory activity of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include assays disclosed in	Miraglia et al., J Biomolecular Screening 4:193-204(1999); Rowland et al., "Lymphocytes: a practical approach" Chapter 6:138-160 (2000); Verhasselt et al., Eur J Immunol 28(11):3886-3890 (1198); Dahlen et al., J Immunol 160(7):3585-3593 (1998); Verhasselt et al., J Immunol 158:2919-2925 (1997); and Nardelli et al., J Leukoc Biol 65:822-828 (1999), the contents of each of which are herein incorporated by reference in its entirety. Human dendritic cells that may be used according to these assays may be isolated using techniques disclosed herein
alpha by dendritic cells			

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S31 Ac Groeler cells	Activation of transcription through the EGR (Early Growth Response) element in immune cells (such as B-cells).	or otherwise known in the art. Human dendritic cells are antigen presenting cells in suspension culture, which, when activated by antigen and/or cytokines, initiate and upregulate T cell proliferation and functional activities.  Assays for the activation of transcription through the EGR response element are well-known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) to regulate EGR transcription factors and modulate expression of immunomodulatory genes. Exemplary assays for transcription through the EGR response element that may be used or routinely modified to test EGR response element activity of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include assays disclosed in: Richards JD, et al., J Immunol, 166(6):3855-3864 (2001); Dinkel, A, et al., J Exp Med, 188(12):2215-2224 (1998); and, Newton, JS, et al., Eur J Immunol 1996 Apr;26(4):811-816 (1996), the contents of each of which are herein incorporated by reference in its entirety. Immune cells that may be used according to these assays include the Raii B-cell line.	additional preferred indication is infection (e.g., an infectious disease as described below under "Infectious Disease").  Preferred embodiments of the invention include using polypeptides of the invention (or antibodies, agonists, or antagonists thereof) in detection, diagnosis, prevention, and/or treatment of Cancer, Autoimmunity, Allergy and Asthma.
Act	hguc	Assays for the activation of transcription through the Gamma Interferon Activation	Preferred embodiments of the invention include using polypeptides of the invention (or antibodies, agonists, or
GA	GAS response element	Site (GAS) response element are well-	antagonists thereof) in detection, diagnosis, prevention,

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and/or treatment of Inflammation, Infection, Cancer, Hypersensitivity, and Atherosclerosis.	A highly preferred indication is diabetes mellitus. An additional highly preferred indication is a complication associated with diabetes (e.g., diabetic retinopathy, diabetic nephropathy, kidney disease (e.g., renal failure,
known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) to regulate STAT transcription factors and modulate gene expression involved in a wide variety of cell functions. Exemplary assays for transcription through the GAS response element that may be used or routinely modified to test GAS-response element activity of polypeptides of the invention (including antibodies and agonists or antagonists of the invention) include assays disclosed in: Gustafson KS, et al., J Biol Chem, 271(33):20035-20046 (1996); Eilers A, et al., Immunobiology, 193(2-4):328-333 (1995); Berger et al., Gene 66:1-10 (1998); Cullen and Malm, Methods in Enzymol 216:362-368 (1992); Henthorn et al., J Immunol 155(10):4582-4587 (1995), the contents of each of which are herein incorporated by reference in its entirety. Exemplary immune cells that may be used according to these assays are publicly available (e.g., through the ATCC). Exemplary immune cells that may be used according to these assays include the U937 cell line, which is a	monocytic cell line. Assays for measuring secretion of insulin are well-known in the art and may be used or routinely modified to assess the ability of polypeptides of the invention (including
in immune cells (such as monocytes).	Stimulation of insulin secretion from pancreatic beta cells.
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